

This Page Is Inserted by IFW Operations  
and is not a part of the Official Record

## **BEST AVAILABLE IMAGES**

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images may include (but are not limited to):

- BLACK BORDERS
- TEXT CUT OFF AT TOP, BOTTOM OR SIDES
- FADED TEXT
- ILLEGIBLE TEXT
- SKEWED/SLANTED IMAGES
- COLORED PHOTOS
- BLACK OR VERY BLACK AND WHITE DARK PHOTOS
- GRAY SCALE DOCUMENTS

**IMAGES ARE BEST AVAILABLE COPY.**

**As rescanning documents *will not* correct images,  
please do not report the images to the  
Image Problems Mailbox.**

**PCT**WORLD INTELLECTUAL PROPERTY ORGANIZATION  
International Bureau

## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification <sup>6</sup> :</b> <b>C12N 15/86, 15/87, 5/10, A61K 35/76, 48/00</b>		<b>A2</b>	<b>(11) International Publication Number:</b> <b>WO 96/39530</b> <b>(43) International Publication Date:</b> 12 December 1996 (12.12.96)
<b>(21) International Application Number:</b> PCT/US96/10245 <b>(22) International Filing Date:</b> 4 June 1996 (04.06.96) <b>(30) Priority Data:</b> 08/462,014 5 June 1995 (05.06.95) US 08/549,489 27 October 1995 (27.10.95) US <b>(60) Parent Applications or Grants</b> <b>(63) Related by Continuation</b> US 08/462,014 (CIP) Filed on 5 June 1995 (05.06.95) US 08/549,489 (CIP) Filed on 27 October 1995 (27.10.95) <b>(71) Applicant (for all designated States except US):</b> THE TRUSTEES OF THE UNIVERSITY OF PENNSYLVANIA [US/US]; 133 South 36th Street, Philadelphia, PA 19104-3246 (US). <b>(72) Inventors; and</b> <b>(75) Inventors/Applicants (for US only):</b> WILSON, James, M. [US/US]; 1350 N. Avignon Drive, Gladwyne, PA 19035 (US). FISHER, Krishna, J. [US/US]; 4006 Pine Street, Philadelphia, PA 19104 (US). GAO, Guang-Ping [CN/US];		<b>Robindale Apartment F5, 1925 Lawrence Road, Havertown, PA 19083 (US).</b> <b>(74) Agents:</b> BAK, Mary, E. et al.; Howson and Howson, Spring House Corporate Center, P.O. Box 457, Spring House, PA 19477 (US). <b>(81) Designated States:</b> AL, AM, AU, AZ, BB, BG, BR, BY, CA, CN, CZ, EE, FI, GE, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LK, LR, LS, LT, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, RO, RU, SD, SG, SI, SK, TJ, TM, TR, TT, UA, UG, US, UZ, VN, ARIPO patent (KE, LS, MW, SD, SZ, UG), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).  <b>Published</b> <i>Without international search report and to be republished upon receipt of that report.</i>	
<b>(54) Title:</b> RECOMBIANT ADENOVIRUS AND ADENO-ASSOCIATED VIRUS, CELL LINES, AND METHODS OF PRODUCTION AND USE THEREOF			
<b>(57) Abstract</b>  An adenovirus E1/E4 expressing packaging cell line is provided, which permits the generation of recombinant adenoviruses deleted in both gene regions. A method for enhancing the efficiency of transduction of a recombinant AAV into a target cell is provided by infecting a target cell with a recombinant AAV comprising a selected transgene under the control of regulatory sequences. The infected cell is contacted with an agent which facilitates the conversion of single stranded recombinant virus to its double stranded form.			

**FOR THE PURPOSES OF INFORMATION ONLY**

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

AM	Armenia	GB	United Kingdom	MW	Malawi
AT	Austria	GE	Georgia	MX	Mexico
AU	Australia	GN	Guinea	NE	Niger
BB	Barbados	GR	Greece	NL	Netherlands
BE	Belgium	HU	Hungary	NO	Norway
BF	Burkina Faso	IE	Ireland	NZ	New Zealand
BG	Bulgaria	IT	Italy	PL	Poland
BJ	Benin	JP	Japan	PT	Portugal
BR	Brazil	KE	Kenya	RO	Romania
BY	Belarus	KG	Kyrgyzstan	RU	Russian Federation
CA	Canada	KP	Democratic People's Republic of Korea	SD	Sudan
CF	Central African Republic	KR	Republic of Korea	SE	Sweden
CG	Congo	KZ	Kazakhstan	SG	Singapore
CH	Switzerland	LI	Liechtenstein	SI	Slovenia
CI	Côte d'Ivoire	LK	Sri Lanka	SK	Slovakia
CM	Cameroon	LR	Liberia	SN	Senegal
CN	China	LT	Lithuania	SZ	Swaziland
CS	Czechoslovakia	LU	Luxembourg	TD	Chad
CZ	Czech Republic	LV	Latvia	TG	Togo
DE	Germany	MC	Monaco	TJ	Tajikistan
DK	Denmark	MD	Republic of Moldova	TT	Trinidad and Tobago
EE	Estonia	MG	Madagascar	UA	Ukraine
ES	Spain	ML	Mali	UG	Uganda
FI	Finland	MN	Mongolia	US	United States of America
FR	France	MR	Mauritania	UZ	Uzbekistan
GA	Gabon			VN	Viet Nam

RECOMBINANT ADENOVIRUS AND ADENO-ASSOCIATED VIRUS,  
CELL LINES, AND METHODS OF PRODUCTION AND USE THEREOF

This invention was supported by the National  
Institute of Health Grant Nos. HD32649-01, DK47757 and  
5 DK49136. The United States government has rights in this  
invention.

Field of the Invention

The present invention relates generally to the field  
of somatic gene therapy, and specifically to methods and  
10 compositions useful in the treatment of genetic  
disorders.

Background of the Invention

Adenoviruses are eukaryotic DNA viruses that can be  
modified to efficiently deliver a therapeutic or reporter  
15 transgene to a variety of cell types [see, e.g., M. S.  
Horwitz et al, "Adenoviridae and Their Replication",  
Virology, second edition, pp. 1712, ed. B. N. Fields et  
al, Raven Press Ltd., New York (1990)]. Recombinant  
adenoviruses (rAds) are capable of providing extremely  
20 high levels of transgene delivery to virtually all cell  
types, regardless of the mitotic state. The efficacy of  
this system in delivering a therapeutic transgene *in vivo*  
that complements a genetic imbalance has been  
demonstrated in animal models of various disorders [K. F.  
25 Kozarsky et al, Somatic Cell Mol. Genet., 19:449-458  
(1993) ("Kozarsky et al I"); K. F. Kozarsky et al, J.  
Biol. Chem., 269:13695-13702 (1994) ("Kozarsky et al II")  
and others]. The use of recombinant adenoviruses in the  
transduction of genes into hepatocytes *in vivo* has  
30 previously been demonstrated in rodents and rabbits [see,  
e.g., Kozarsky II, cited above, and S. Ishibashi et al,  
J. Clin. Invest., 92:883-893 (1993)].

The first-generation recombinant, replication-  
deficient adenoviruses which have been developed for gene  
35 therapy contain deletions of the entire E1a and part of  
the E1b regions. This replication-defective virus is

grown on an adenovirus-transformed, complementation human embryonic kidney cell line containing a functional adenovirus E1a gene which provides a transacting E1a protein, the 293 cell [ATCC CRL1573]. E1-deleted viruses are capable of replicating and producing infectious virus in the 293 cells, which provide E1a and E1b region gene products in trans. The resulting virus is capable of infecting many cell types and can express the introduced gene (providing it carries its own promoter), but cannot replicate in a cell that does not carry the E1 region DNA unless the cell is infected at a very high multiplicity of infection.

Adeno-associated virus (AAV) is an integrating human DNA parvovirus which has been proposed for use as a gene delivery vehicle for somatic gene therapy [B. J. Carter, in "Handbook of Parvoviruses", ed., P. Tijsser, CRC Press, pp.155-168 (1990)]. This small non-enveloped virus contains a 4.6 kb single stranded (ss) DNA genome that encodes sets of regulatory and capsid genes called rep and cap. Rep polypeptides (rep78, rep68, rep62 and rep40) are involved in replication, rescue and integration of the AAV genome. The cap proteins (VP1, VP2 and VP3) form the virion capsid. Flanking the rep and cap open reading frames at the 5' and 3' ends are 145 bp inverted terminal repeats (ITRs), the first 125 bp of which are capable of forming Y- or T-shaped duplex structures.

Recombinant forms of AAV (rAAV) have been developed as vectors by replacing all viral open reading frames with a therapeutic minigene, while retaining the necessary cis elements contained in the ITRs. [See, e.g., US Patent Nos. 4,797,368; 5,153,414; 5,139,941; 5,252,479; and 5,354,678; and International Publication Nos. WO 91/18088 published November 28, 1991; WO 93/24641 published December 9, 1993 and WO94/13788 published June

23, 1994]. However, progress towards establishing AAV as a transducing vehicle for gene therapy has been slow for a variety of reasons. For example, the integrated provirus preferentially targets specific sites in chromosome 19. Additionally, difficulties surround large-scale production of replication defective recombinants. The cells employed to produce rAAV must also be infected with adenovirus or herpesvirus to provide the necessary helper functions, thereby producing problems in purifying recombinant AAV (rAAV) from contaminating virus in culture. Practical experience with purified recombinant AAV as a gene therapy vector has been disappointing, because the more purified the AAV is from co-infection with its helper virus in culture, the lower the gene transduction efficiencies that the rAAV displays.

There remains a need in the art for additional recombinant adenoviruses and rAAV, therapeutic compositions and methods which enable effective use of these recombinant viruses in the treatment of disorders and diseases by gene therapy.

#### Summary of the Invention

In one aspect of this invention, a packaging cell line is provided which expresses adenovirus genes E1a, E1b and E4, or functional fragments thereof, e.g., the E4 open reading frame (ORF) 6.

In another aspect, the invention provides a rAd comprising the DNA of at least a portion of the genome of an adenovirus having functional deletions of the E1 and E4 gene regions; a suitable gene operatively linked to regulatory sequences directing its expression, and an adenovirus capsid, the rAd capable of infecting a mammalian cell and expressing the gene product in the cell *in vivo* or *in vitro*. The invention also provides a mammalian cell infected with the rAd described above.

In still another aspect, the invention provides a rAd shuttle vector comprising the DNA of at least a portion of the genome of an adenovirus having functional deletions of the E1 and E4 gene regions.

5 In a further aspect, the invention provides a method for producing the above-described recombinant Ad and a method for delivering a selected gene into a mammalian cell using the recombinant Ad described above.

10 In another aspect, the invention provides a method for enhancing the efficiency of transduction of a recombinant AAV into a target cell. The method operates, in brief, by infecting a target cell with a ss recombinant adeno-associated virus (rAAV) which comprises a transgene operatively linked to regulatory sequences  
15 directing its expression, and contacting the infected cells with an agent which facilitates the conversion of ss rAAV to its double stranded (ds) form. Conversion of ss rAAV to ds rAAV occurs in the target cell, resulting in enhanced transduction of the rAAV into the target  
20 cell. The agent may be a helper virus which carries a selected gene or functional fragment thereof encoding a polypeptide capable of enhancing the conversion of the ss rAAV to ds rAAV and which is co-infected into the same target cell. The agent may also be a drug or chemical  
25 composition which accomplishes the same function and is applied to the infected target cell. This method can operate both in an ex vivo setting and in vivo.

In yet another aspect, the invention provides a novel recombinant AAV, which contains both the transgene  
30 intended for use in treating a genetic disease or disorder and at least one additional gene operatively linked to inducible or constitutive regulatory sequences. The additional gene(s) encodes a polypeptide capable of facilitating, alone or in concert with other additional  
35 genes, the conversion of ss rAAV to its ds form upon

expression. In a preferred embodiment, the additional gene is adenovirus E4 or a functional fragment thereof. Also disclosed is a method for enhancing the efficiency of transduction of the novel rAAV into a target cell.

5 The novel rAAV and methods of this invention are also useful in pharmaceutical compositions for use in *ex vivo* and *in vivo* gene therapy treatment protocols for treating inherited diseases, cancer, and other genetic dysfunctions.

10 Other aspects and advantages of the present invention are described further in the following detailed description of the preferred embodiments thereof.

#### Brief Description of the Drawings

Fig. 1 is a schematic drawing of an exemplary  
15 plasmid pMMTVE4ORF6 [SEQ ID NO: 1] or pMTE4ORF6, which contains an MMTV or sheep MT promoter, respectively, in control of a human E4 ORF 6 gene sequence, a growth hormone gene terminator sequence (GH), an SV40 ori, pBR322-based plasmid sequences including a neo<sup>R</sup> gene, an  
20 SV40 polyA site and an amp<sup>R</sup> gene.

Fig. 2 is a schematic map of rAd H5.001CBLacZ [SEQ ID NO: 3] with indicated restriction endonuclease enzyme sites. The striated bar represents the CBLacZ minigene; the black bar represents Ad5 viral backbone, the  
25 crosshatched bar represents Ad E4 deletion.

Fig. 3 plots LacZ forming units (LFU)/ml vs time (hours) for E4 complementing cell lines infected with H5.001CBLacZ.

Fig. 4A is a graph of the induction, ORF6 expression  
30 and viral production in 293-27-18 packaging cells plotting yield at 24 hours post-infection (pi) in LFU/ml and ORF6 protein (abs.mm) vs. concentration of the inducer, dexamethasone ( $\mu$ M). Abs.mm is the intensity of the size of the protein band on a Western blot and  
35 reflects absorbance and protein size in mm<sup>2</sup>. The square



is yield at 24 hours pi. The diamond is ORF6 prot in detected at 24 hours pi.

Fig. 4B is a similar graph to that of Fig. 4A, except that the packaging cells are 293-10-3 cells. The symbols are as described for Fig. 4A.

Fig. 5A is a bar graph plotting  $\beta$ -galactosidase enzyme activity in lysates from infected HeLa cells. The horizontal axis indicates the adenoviruses infected into the HeLa cells, with the symbol "+" indicating the addition of the adenovirus to the rAAV, AV.CMVLacZ. The vertical axis indicates intracellular  $\beta$ -galactosidase specific activity (mUnits/mg protein) using ONPG. Below each bar, the fold-induction in specific activity relative to cells that received the AV.CMVLacZ vector alone is given.

Fig. 5B is a bar graph plotting Ad multiplicity of infection (MOI) in HeLa cells of wild-type Ad5 or the E2 mutant dl802, the cells co-infected with rAAV vs. intracellular  $\beta$ -galactosidase specific activity. See Example 11.

Fig. 6A is a graph in which  $\beta$ -galactosidase specific activity and counts per minute (CPM) are plotted along the vertical axis and adenovirus MOI's are on the horizontal axis for HeLa cells infected with wtAd5 and rAAV according to Example 12. Data obtained from low MOI (1, 5, and 10) infections are shown.

Fig. 6B is a graph similar to that of Fig. 6A except that the cells were infected with Ad mutant dl802.

Fig. 7A illustrates a model for leading strand synthesis of a complementary AAV strand in the presence of Rep (+Rep) or absence of Rep (-Rep). Rep expresses a terminal resolution activity that can convert a duplex structure with closed-ends to an open-ended duplex. In the absence of Rep, terminal resolution is impaired leaving the covalently closed, hairpin structures intact.

Under these conditions, hairpins are expected to be found leftward and rightward, since both strands of a rescued ds AAV genome are packaged into virions.

Fig. 7B is a schematic of linear AV.CMV $\text{LacZ}$  with  
5 labeled domains including the AAV ITRs, CMV immediate early enhancer/promoter (CMV), SV40 splice donor-splice acceptor (SD/SA), *E. coli*  $\beta$ -galactosidase cDNA ( $\text{LacZ}$ ), and SV40 polyA signal (pA). Two NotI sites located at bp positions 1035 and 4509 are indicated.

10 Fig. 7C illustrates a closed end and an open end fragment of rAV.CMV $\text{LacZ}$ .

Figs. 7D, 7E and 7F indicate the mixture of open-ended and covalently closed duplex fragments generated by NotI digestion of ss AV.CMV $\text{LacZ}$  at position  
15 4509 in the absence of terminal resolution. The NotI 4509 digestion provides a convenient means of releasing a 361 bp fragment that contains the right ITR in the context of a hybridization target (i.e. SV40 pA). In the presence of terminal resolution, only the open-ended 361  
20 bp fragment would be expected to be generated (Fig. 7D) by such digestion.

Fig. 8A is a bar graph plotting  $\beta$ -galactosidase specific activity (mUnits/mg protein) vs. increasing concentration of zinc ( $\mu\text{M}$ ) inducer for cell line 293 (MT-  
25 ORF6) transduced with AVCMV $\text{LacZ}$  (first row below each bar). Also provided is the fold-induction relative to 293 cells (second row below each bar), and the fold-induction relative to 293(ORF6) cells maintained in the absence of zinc (third row).

30 Fig. 8B is a bar graph plotting CPM of duplex monomer replicative form (RFm) of rAAV vs. the concentration of zinc ( $\mu\text{M}$ ) used for induction and the fold-induction relative to 293(ORF6) cells maintained in 0 mM zinc below each bar.

Fig. 8C is a graphical comparison of the induction profiles that describe AV.CMV $\text{LacZ}$  transduction efficiency. Specific activity data from Fig. 8A and CPM data of AV.CMV $\text{LacZ}$  RFm from Fig. 8B are plotted along the vertical axis, and concentration of zinc sulfate used during the experiment is shown along the horizontal axis.

Fig. 9 is a bar graph plotting specific activity (milliunits  $\beta$ -galactosidase/mg protein) vs the concentration of zinc used for induction (first row under the horizontal axis), the fold-induction relative to HeLa cells (second row), and the fold-induction relative to HeLa(Mt-ORF6) cells maintained in the absence of zinc (third row), for the HeLa(MT-ORF6) cells transduced at an MOI of 1,000 AV.CMV $\text{LacZ}$  virus particles/cell in the absence of zinc sulfate inducer or in the presence of 50, 100, 150, 200 or 250  $\mu\text{M}$  zinc sulfate inducer.

Fig. 10 is a schematic of the plasmid pAV.CMV $\text{LacZ}$  [SEQ ID NO: 4].

Fig. 11 illustrates plasmid pAV.CMVALP.GRE-ORF6 [SEQ ID NO: 5].

#### Detailed Description of the Invention

The present invention provides packaging cell lines, which enable the production of recombinant adenoviruses (rAd) functionally deleted in both the E1 and E4 genes. These rAd and methods which enable the therapeutic treatment of disorders with such rAds are disclosed. Novel "second generation" recombinant adeno-associated virus (rAAV) and methods for enhancing the transduction efficiency of rAAV containing a transgene for expression in a somatic gene therapy protocol are also provided. The methods and compositions of this invention are useful in ex vivo applications of gene therapy, such as in the transduction of bone marrow cells with desirable hematopoietic stem cell progenitor genes prior to bone marrow transplantation. The embodiments of the invention

are also useful in pharmaceutical compositions for direct *in vivo* treatment of patients by gene therapy vectors, including the transduction of desirable genes in patients with genetic disorders, such as cystic fibrosis.

5           I. *Packaging Cell Lines*

To increase the transgene capacity and decrease immune response of rAds, as many viral genes as possible should be deleted to inactivate the adenovirus. However, it is crucial to generate complementing cell lines for construction and propagation of such deleted Ad. The method and compositions of the present invention overcome several problems previously identified in the gene therapy for first generation E1 deleted adenoviruses and display advantages in administration particularly to muscle tissue.

Early region 4 (E4) of Ad serotype 5 consists of 7 ORFs believed to be involved in viral DNA replication, host cell shut-off, and late mRNA accumulation. To generate rAd deleted in E4, the function of the E4 region must be supplied to the rAd by a helper virus or packaging cell line. However, useful packaging cell lines have not been available previously because normally the continuous expression of functioning Ad E1 and functional E4 in a single cell line are toxic to the cell. Such cells are therefore not useful for the growth and replication of rAds. Further, the DNA encoding the functional Ad E1 and Ad E4 genes, when present in a packaging cell line, can increase the chances of recombination with a rAd virus to cause the virus to revert to a wildtype Ad virus.

The present invention avoids these problems by providing a packaging cell line which contains the Ad5 E1 gene and only the ORF 6 of the Ad5 E4 gene. ORF6 of E4 alone can provide the requirements for E4 in the viral life cycle.

According to this invention, the ORF6 is preferably under the transcriptional control of an inducible promoter. The mouse mammary tumor virus (MMTV) promoter, inducible by a glucocorticoid, particularly, dexamethasone, is presently preferred. The DNA sequence of the MMTV promoter spans nucleotides 1-1506 of SEQ ID NO: 1. Another inducible promoter is the sheep metallothionine (MT) promoter, inducible by zinc [M. G. Peterson et al, Eur. J. Biochem., 174:417-424 (1988)]. However, the zinc sulfate inducer of the MT promoter can itself be toxic to the cells. Other inducible promoters, such as those identified in International patent application WO95/13392, published May 18, 1995, and incorporated by reference herein may also be used in the production of packaging cell lines according to this invention. Constitutive promoters, such as the constitutive Ad5 E4 region promoter, LTR, may be employed in control of the expression of ORF6.

The packaging cell line of the invention which utilizes an inducible promoter permits one to control the development of toxicity by regulating the expression of the E4 ORF6 gene. After the desired shuttle vector containing the Ad sequences is transfected into the cell line, expression of the E4 ORF6 can be induced by the appropriate inducer. The packaging cell is thus able to provide both Ad E1 and Ad E4 ORF6 gene products to the rAd for a sufficient period to allow productive infection and recovery of the rAd, before the cell becomes toxic. At present, the time period before the cell experiences toxicity is about 10 days.

In its most preferred form, the packaging cell line is a human embryonic kidney (HEK) 293 E1 expressing cell line into which is introduced the E4 ORF 6 sequence under the control of the inducible promoter. It should be understood by one of skill in the art that another parent

cell line may be selected for the generation of a novel cell line expressing the E1a, E1b, and E4 ORF6 genes of a selected adenovirus serotype. Among such parent cell lines may be included HeLa [CCL 2], A549 [CCL 185], KB [CCL 17], Detroit [e.g., Detroit 510, CCL 72] and WI-38 [ATCC CCL 75] cells. These cell lines are all available from the American Type Culture Collection, 12301 Parklawn Drive, Rockville, MD, USA. Other suitable parent cell lines may be obtained from other sources. If such parent cell lines were selected for modification, the cell line would need to be further supplied with the E1a and E1b gene functions, e.g., such as by transfection with a plasmid containing these genes or functional fragments thereof under a suitable promoter, as well as with the ORF6 gene as described herein.

Example 1 teaches construction of packaging cell lines containing only the ORF 6 of Ad5 E4 region or, for functional comparisons, the entire E4 region. Briefly described, the entire E4 region and an ORF6 sequence of Ad 5 E4 gene were obtained by known techniques [see, e.g., Sambrook et al., "Molecular Cloning. A Laboratory Manual.", 2d edit., Cold Spring Harbor Laboratory, New York (1989) and references cited therein]. To isolate the ORF6 region, the anchored PCR technique was used to amplify the ORF6 sequence from its initiation codon to its termination codon. Primers selected from the published sequence of ORF6 were used to amplify the ORF sequence and insert restriction sites onto the end of the sequence. The E4 ORF6 sequence itself is reproduced as nucleotides 1523 through 2408 of SEQ ID NO: 1. The entire E4 gene sequence is published in the Genbank sequence of Ad5 [Genbank Accession No. M73260].

A minigene was constructed that placed the ORF6 sequence under the control of a selected promoter. By "minigene" as used here is meant the combination of the

desired sequence to be expressed (in this particular instance, the ORF6 sequence) and the other regulatory elements necessary to transcribe the desired sequence and express the gene product in a cell containing that minigene. The ORF6 sequence gene is operatively linked to regulatory components in a manner which permits its transcription. Such components include conventional regulatory elements, such as a promoter to drive ORF6 expression. One inducible promoter was the  $Zn^{+2}$  inducible MT promoter; the other was the dexamethasone-inducible MMTV promoter of SEQ ID NO: 1.

The minigene also contains nucleic acid sequences heterologous to the ORF6 viral sequence, including sequences providing signals required for efficient polyadenylation of the transcript (poly-A or pA). A common poly-A sequence which is employed in this invention is that derived from the growth hormone (GH) gene terminator sequence (nuc. 2409-3654 of SEQ ID NO: 1). The poly-A sequence generally is inserted in the minigene following the ORF6 sequence. The polyA sequence employed in the MMTV-ORF6 minigene described in Example 1 [SEQ ID NO: 1] is supplied by the GH gene terminator and an SV40 origin of replication (ori). A similar minigene differing in promoter sequence, polyA sequence and/or SV40 ori can also be designed by one of skill in the art to transfer the E4 ORF6 sequence to a shuttle plasmid. Selection of these and other common vector elements are conventional [see, e.g., Sambrook et al, cited above, and references cited therein] and many such sequences are available from commercial and industrial sources as well as from Genbank.

The ORF6-containing minigene was subcloned into a pBR322-based shuttle plasmid that contained a neomycin resistance gene, resulting in the shuttle vector of Fig. 1. Any of the many known bacterial shuttle vectors may

be employed to carry the minigene, providing that the vector contains a reporter gene or selectable marker of which many, e.g., neo, amp or purimycin, are known in the art. It is expected that one of skill in the art can  
5 develop other suitable shuttle vectors using other plasmid components which are similarly capable of transferring the ORF6 minigene into the chromosome of a cell transfected with the plasmid.

As further described in Example 1, other shuttle  
10 vectors were designed for comparative purposes, which contain the complete or substantially complete Ad5 E4 region under the control of the constitutive retroviral MLV LTR sequence in the presence or absence of the endogenous E4 promoter. The shuttle plasmid carrying the  
15 ORF6 minigene (or the entire E4 region) was introduced into HEK 293 cells which express the Ad E1 gene products. Complementing cell lines were generated that express these Ad E4 or ORF6 genes from either their endogenous promoters or heterologous inducible promoters. These  
20 cell lines are further characterized by their genetic constitution, E4 protein synthesis, recombinant AAV helper function, relative plaque efficiency of H5dl1004 virus, and growth kinetics of recombinant E1/E4 deleted adenovirus. These characteristics of exemplary E1/E4  
25 expressing packaging cell lines are discussed in detail in the following examples.

## *II. Recombinant Adenovirus*

The E1/E4 expressing cell line is useful in constructing E1/E4 deleted rAds which can deliver a  
30 suitable gene to mammalian cells and tissues. These rAd are functionally deleted in at least the E1a, E1b and E4 Ad gene regions. By the term "functionally deleted" is meant that a sufficient amount of the gene region is removed or otherwise damaged, e.g., by mutation or  
35 modification, so that the gene region is no longer



capable of producing the products of gene expression. If desired, the entire gene region may be removed. In *in vivo* experiments with the rAd grown in the packaging cell lines, the E1/E4 deleted rAd demonstrated utility particularly in transferring a transgene to a muscle cell.

The adenovirus sequences used in the construction of the shuttle vectors, helper viruses, if needed, and rAd, and other components and sequences employed in the construction of the vectors and viruses described herein may be readily obtained from commercial or academic sources based on previously published and described sequences. Viral materials may also be obtained from an individual patient. The viral sequences and vector components may be generated by resort to the teachings and references contained herein, coupled with standard recombinant molecular cloning techniques known and practiced by those skilled in the art. Modifications of existing nucleic acid sequences forming the vectors, including sequence deletions, insertions, and other mutations taught by this specification may be generated using standard techniques. Similarly, the methods employed for the selection of viral sequences useful in a vector, the cloning and construction of the "minigene" and its insertion into a desired viral shuttle vector and the production of a recombinant infectious virus are within the skill in the art given the teachings provided herein.

*A. Construction of the Transgene*

A "minigene" in this context is defined as above, except that the components of this minigene are designed to express the gene product *ex vivo* or *in vivo*. Such components include conventional regulatory elements necessary to drive expression of the transgene in a cell transfected with the rAd. For this minigene, a selected

promoter is operatively linked to the transgene and located, with other regulatory elements, within the selected viral sequences of the recombinant vector. Selection of the promoter is a routine matter and is not  
5 a limitation of this invention. Useful promoters may be constitutive promoters or regulated (inducible) promoters, which will enable control of the amount of the transgene to be expressed. For example, a desirable promoter is that of the cytomegalovirus (CMV) immediate  
10 early promoter/enhancer [see, e.g., Boshart et al, Cell, 41:521-530 (1985)]. Another desirable promoter includes the Rous sarcoma virus LTR promoter/enhancer. Still another promoter/enhancer sequence is the chicken cytoplasmic  $\beta$ -actin (CB) promoter [T. A. Kost et al,  
15 Nucl. Acids Res., 11(23):8287 (1983)]. Other suitable promoters may be selected by one of skill in the art.

The minigene may also desirably contain nucleic acid sequences heterologous to the viral vector sequences including poly-A sequences and introns with functional  
20 splice donor and acceptor sites, as described above. The poly-A sequence generally is inserted in the minigene following the transgene sequences and before the 3' adenovirus sequences. A minigene of the present invention may also contain an intron, desirably located  
25 between the promoter/enhancer sequence and the transgene. Selection of these and other common vector elements are conventional as described above and many such sequences are available from commercial and industrial sources as well as from Genbank.

30 As above stated, the minigene is located in the site of any selected deletion in the rAd. In the E1/E4 deleted rAd H5.001CBLacZ, the transgene is located in the deleted E1 gene region. However, the transgene may be located elsewhere in the adenovirus sequence, as desired.

*B. Production of Recombinant Adenovirus*

Adenovirus sequences useful in this invention may include the DNA sequences of a number of adenovirus types, which are available from Genbank, including type Ad5 [Genbank Accession No. M73260]. The adenovirus sequences may be obtained from any known adenovirus serotype, such as serotypes 2, 3, 4, 7, 12 and 40, and further including any of the presently identified 41 human types [see, e.g., Horwitz, cited above]. Similarly adenoviruses known to infect other animals may also be employed in the vector constructs of this invention. The selection of the adenovirus type is not anticipated to limit the following invention. A variety of adenovirus strains are available from the American Type Culture Collection, Rockville, Maryland, or available by request from a variety of commercial and institutional sources. In the following exemplary embodiment an adenovirus, type 5 (Ad5) is used for convenience.

However, it is desirable to obtain a variety of adenovirus shuttle vectors based on different human adenovirus serotypes. It is anticipated that a library of such plasmids and the resulting rAds would be useful in a therapeutic regimen to evade cellular, and possibly humoral, immunity, and lengthen the duration of transgene expression, as well as improve the success of repeat therapeutic treatments. Additionally the use of various serotypes is believed to produce rAd with different tissue targeting specificities. Additionally, the absence of adenoviral genes E1 and E4 in the rAd of this invention should reduce or eliminate adverse CTL responses which normally cause destruction of rAds deleted of only the E1 gene.

rAds of this invention are recombinant, defective adenoviruses (i.e., E1 deleted) which are also deleted completely or functionally of the E4 gene region.

Functional deletions of E4 gene regions may be assessed by assays of Examples 2 and 3, among other assays. rAds useful in this invention may optionally bear other mutations, e.g., temperature sensitive (ts) mutations in the E2a gene region, and deletions in the E3 gene regions.

An adenovirus of this invention contains a functional deletion of the adenoviral early immediate early gene E1a (which spans mu 1.3 to 4.5) and delayed early gene E1b (which spans mu 4.6 to 11.2). Similarly the adenovirus has a functional deletion of the whole E4 region (which spans mu 92 to 97.2), or of at least ORF6 of the E4 region. Gene regions which may be optionally deleted in the E1/E4 deleted rAd of this invention include all or a portion of the adenovirus delayed early gene E3 (which spans mu 76.6 to 86.2). The function of E3 is irrelevant to the function and production of the rAd.

The rAd of this invention may also have a mutation which results in reduced expression of adenoviral protein and/or reduced viral replication. For example, a ts mutation may be introduced into the adenovirus delayed early gene E2a (which spans mu 67.9 to 61.5). Among such mutations include the incorporation of the missense ts mutation in the (DBP)E2a region found in the Ad5 H5ts125 strain [P. Vander Vliet et al, J. Virol., 15:348-354 (1975)] at 62.5 mu. A single amino acid substitution (62.5 mu) at the carboxy end of the 72 kd protein produced from the E2a gene in this strain produces a protein product which is a ss DNA binding protein and is involved in the replication of adenoviral genomic DNA. At permissive temperatures (approximately 32°C) the ts strain is capable of full life cycle growth on HeLa cells, while at non-permissive temperatures (approximately 38°C) no replication of adenoviral DNA is

seen. In addition, at non-permissive temperatures, decreased immunoreactive 72 kd protein is seen in HeLa cells. See, e.g., J. F. Engelhardt et al, Hum. Gene Ther., 5:1217-1229 (1994); J. F. Engelhardt et al, Proc. Natl. Acad. Sci., USA, 91:6196-6200 (1994) and International patent application WO95/13392, published May 18, 1995, incorporated by reference herein.

However, it should be understood that other deletions in the adenovirus genome as previously described in the art or otherwise may also occur in the rAds of this invention. One minimal type of rAd can contain adenovirus genomic sequences from which all viral genes are deleted. More specifically, the adenovirus sequences may be only the cis-acting 5' and 3' inverted terminal repeat (ITR) sequences of an adenovirus (which function as oris) and the native 5' packaging/enhancer domain, that contains sequences necessary for packaging linear Ad genomes and enhancer elements for the E1 promoter. The adenovirus 5' sequence containing the 5' ITR and packaging/enhancer region (Ad5 mu 0-1 or bp 1-360) can be employed as the 5' adenovirus sequence in rAd of this invention. The 3' adenovirus sequences including the right terminal (3') ITR sequence of the adenoviral genome spanning about bp 35,353 - end of the adenovirus genome, or map units 98.4-100 may be desirably employed as the 3' sequence of the rAd. These sequences, which are clearly devoid of the E1 and E4 genes, can flank, or be operatively associated with the minigene in a rAd. Any other necessary Ad gene products will then be supplied by helper viruses and the E1/E4 ORF6 expressing packaging cell of this invention.

Exemplary rAd for use in this invention, for example, may be obtained by homologous recombination of desired fragments from various rAd, a technique which has been commonly employed to generate other rAd for gene

therapy use. In the examples below, a representative rAd, H5.001CBLacZ, is constructed by homologous recombination between the adenovirus dl1004 (also H5dl1004) viral backbone and pAdCBLacZ minigene DNA.

- 5 H5dl1004 is an Ad5 virus deleted of from about map unit 92.1 through map unit 98, i.e, substantially the entire E4 gene. The dl1004 virus is described in Bridge and Ketner, J. Virol., 632(2):631-638 (Feb. 1989).

- 10 The pAdCBLacZ vector is a cDNA plasmid containing Ad m.u. 0-1, an E1 deletion into which is inserted a bacterial  $\beta$ -galactosidase gene under the control of a chicken  $\beta$ -actin promoter, with other regulatory elements as described below, and flanked by Ad m.u. 9-16 and plasmid sequence.

- 15 The production of the E1/E4 rAd of this invention in the packaging cell line of this invention utilizes conventional techniques. Such techniques include conventional cloning techniques of cDNA such as those described in texts [Sambrook et al, cited above],  
20 use of overlapping oligonucleotide sequences of the adenovirus genomes, PCR and any suitable method which provides the desired nucleotide sequence. Standard transfection and co-transfection techniques are employed, e.g., CaPO<sub>4</sub> transfection techniques using the  
25 complementation 293 cell line. Other conventional methods employed include homologous recombination of the viral genomes, plaquing of viruses in agar overlay, methods of measuring signal generation, and the like.

- For example, following the construction and  
30 assembly of the desired minigene-containing plasmid vector pAdCBLacZ, the E1/E4 expressing packaging cell line of this invention is infected with the helper virus H5dl1004. The infected cell line is then subsequently transfected with an adenovirus plasmid vector by  
35 conventional methods. Homologous recombination occurs

between the E4-d let d H5dl1004 helper and the pAdCBLacZ vector, which permits the adenovirus-transgene sequences in the vector to be replicated and packaged into virion capsids, resulting in the rAd. About 30 or more hours post-transfection, the cells are harvested, an extract prepared and the rAd containing the LacZ transgene is purified by buoyant density ultracentrifugation in a CsCl gradient.

### *III. Use of the Recombinant Virus in Gene Therapy*

The rAd containing the transgene produced by cooperation of the adenovirus vector and E4 deleted helper virus and packaging cell line, as described above, provides an efficient gene transfer vehicle which can deliver the transgene in a pharmaceutical composition to a patient *in vivo* or *ex vivo* and provide for integration of the gene into a mammalian cell.

The rAd are administered to humans in a conventional manner for gene therapy and serve as an alternative or supplemental gene therapy for the disorder to which the transgene is directed. A rAd of this invention may be administered to a patient, preferably suspended in a biologically compatible solution or pharmaceutically acceptable delivery vehicle. A suitable vehicle includes sterile saline. Other aqueous and non-aqueous isotonic sterile injection solutions and aqueous and non-aqueous sterile suspensions known to be pharmaceutically acceptable carriers and well known to those of skill in the art may be employed for this purpose.

The rAd are administered in sufficient amounts to transfect the desired target cells, e.g., muscle, liver, epithelial, etc. and provide sufficient levels of transfer and expression of the transgene to provide a therapeutic benefit without undue adverse or with medically acceptable physiological effects which can be determined by those skilled in the medical arts.

Conventional and pharmaceutically acceptable routes of administration include direct delivery to the muscle or other selected cell, intranasal, intravenous, intramuscular, subcutaneous, intradermal, oral and other parental routes of administration. Routes of administration may be combined, if desired.

Dosages of rAd will depend primarily on factors such as the condition being treated, the age, weight and health of the patient, and may thus vary among patients. For example, a therapeutically effective human dose of the rAd is generally in the range of from about 20 to about 100 ml of saline solution containing concentrations of from about  $1 \times 10^9$  to  $1 \times 10^{11}$  pfu/ml virus. A preferred human dose is estimated to be about 50 ml saline solution at  $2 \times 10^{10}$  pfu/ml. The dose will be adjusted to balance the therapeutic benefit against any side effects. The levels of expression of the transgene can be monitored to determine the frequency of administration.

An optional method step involves the co-administration to the patient, either concurrently with, or before or after administration of the rAd of a suitable amount of a short acting immune modulator. The selected immune modulator is defined herein as an agent capable of inhibiting the formation of neutralizing antibodies directed against the recombinant vector of this invention or capable of inhibiting or substantially delaying cytolytic T lymphocyte (CTL) elimination of the vector. Among desirable immune modulators are interleukin-12 [European Patent Application No. 441,900]; gamma interferon [S. C. Morris et al, J. Immunol., 152:1047 (1994)]; interleukin-4 [United States Patent No. 5,017,691]; antibody to the CD4 protein, such as anti-OKT 3+ [see, e.g., US Patent No. 4,658,019] or antibody GK1.5 (ATCC Accession No. TIB207); a soluble CD40 molecule or an antibody to CD40 ligand (Bristol-Myers Squibb Co)



[European patent application 555,880, published August 18, 1993]; a soluble form of B7 or an antibody to CD28 or CTLA4 [CTLA4-Ig (Bristol-Myers Squibb Co), European patent application 606,217, published July 20, 1994], or agents such as cyclosporin A or cyclophosphamide. Thus, the pharmaceutical compositions and methods of this invention provide a desirable gene therapy treatment.

#### IV. Recombinant Adeno-Associated Virus

In the following context the term "transgene" means a nucleic acid sequence or reverse transcript thereof, heterologous to the AAV sequence, which encodes a polypeptide or protein of interest. The transgene may be operatively linked to regulatory components in a manner which permits transgene transcription, i.e., the transgene is placed into operative association with a promoter, as well as other regulatory sequences, such as SV40 introns or polyA sequences, useful for its regulation. The composite association of the transgene with its regulatory sequences is referred to herein as a minicassette or minigene.

The composition of the transgene or minicassette sequence will depend upon the use to which the resulting rAAV will be put. For example, one type of transgene sequence includes a reporter sequence, which upon expression produces a detectable signal. Such reporter sequences include without limitation, an *E. coli*  $\beta$ -galactosidase (*LacZ*) cDNA, an alkaline phosphatase gene (ALP) and a green fluorescent protein gene. These sequences, when associated with regulatory elements which drive their expression, provide signals detectable by conventional means, e.g., ultraviolet wavelength absorbance, visible color change, etc.

Another type of transgene sequence includes a therapeutic gene which expresses a desired gene product in a host cell. These therapeutic nucleic acid sequences

typically encode products for administration and expression in a patient *in vivo* or *ex vivo* to replace or correct an inherited or non-inherited genetic defect or treat an epigenetic disorder or disease. Such transgenes  
5 may be readily selected by one of skill in this art and the design of the transgene or the minicassette for insertion into the rAAV is not a limitation of this invention.

The term "rAAV" encompasses any recombinant AAV gene therapy vehicle of the prior art, including the AdAAV hybrid virus described in published International Patent Application No. WO96/13598, published May 9, 1996. More specifically, rAAV defines a rAAV comprising: (a) the DNA of at least a portion of the genome of an AAV, which  
15 portion is capable of transducing into a target cell at least one selected gene in the absence of cell division; and (b) at least one selected gene (or transgene) operatively linked to regulatory sequences directing its expression, the gene flanked by the DNA of (a) and  
20 capable of expression in the target cell *in vivo* or *in vitro*.

Other rAAVs have been described in the art. The method of this invention is not limited by the precise nature of the AAV sequences used in the rAAV, provided  
25 that at a minimum both the 5' and 3' AAV inverted terminal repeats are present. Thus, the rAAV may be selected by one of skill in the art, and is not itself a limitation on this invention. The rAAVs specifically disclosed herein are illustrative.

By the term "transduction" is meant that the rAAV produced by practice of the invention is capable of infecting a desired target cell and expressing the transgene in the cell by harnessing the cell's machinery. Transduction may include stably integrating the viral DNA  
30 into a chromosome of the target cell. "Enhanced

transduction" is defined as the ability of the rAAV in the presence of a conversion agent to transduce the target cell, either *in vitro*, *ex vivo* or *in vivo*, at an efficiency greater than a typical prior art rAAV produced in, and purified from, a culture co-infected with an adenovirus or herpesvirus helper.

This method is based on the observation that the limiting step in rAAV mediated transduction of cells for gene therapy is not the internalization or transfer of the ss viral genome, but rather the subsequent conversion of the single-stranded (ss) viral genome to a transcriptionally active double-stranded (ds) form. Formation of ds DNA intermediates is necessary for recombinant gene expression, which is likely to be modulated by viral and cellular factors through posttranscriptional mechanisms. The inventors have designed a method to overcome this rate-limiting step, thereby enhancing transduction ability of an rAAV and ultimately the use of rAAV in gene therapy protocols.

This method of the present invention may employ a conventionally prepared ss rAAV containing a transgene. The prior art produces ss rAAV by co-infection in culture with a helper adenovirus or herpesvirus, followed by purifying the rAAV from the culture contaminants including the helper virus, and infecting the target cell with the rAAV alone. The present invention provides for infecting a target cell with a ss rAAV. However, once the target cell is infected, the infected cell is contacted with an agent which facilitates the conversion of the ss rAAV to the ds form of rAAV. The action of this "facilitating agent" or "conversion agent" causes the ss to ds conversion to occur in the target cell, resulting in enhanced transduction of the recombinant AAV into the target cell. By facilitating the conversion of ss to ds rAAV in the target cell, the method of this

invention may also result in both transduction and stable chromosomal integration of the rAAV into the chromosome of said host cell.

Preferably, for use of this invention the  
5 "facilitating or conversion agent" may take several forms.

*A. The Conversion Agent is a Helper Virus*

In one embodiment, the agent is a helper virus and the method includes an additional step of co-  
10 infecting the target cell with the helper virus. The helper virus useful in this method contains a selected gene which can facilitate the conversion of ss rAAV to ds rAAV. The selected gene may encode a gene product or polypeptide (or a functional fragment of the polypeptide  
15 which shares the biological activity of the full-length polypeptide) which enhances the conversion.

Alternatively, the selected gene may express an antisense or ribozyme which functions in the cell to block or  
20 inhibit a cellular gene that normally prevents ss to ds conversion of the rAAV. These genes may also be employed in the second generation rAAV described below.

The helper virus is capable of expressing the selected gene product in the target cell in the absence of cell division. The helper virus may be a wild-type or  
25 mutant adenovirus. The helper virus may alternatively be a wild-type or mutant herpesvirus. Preferably, for use as facilitating agents, such viruses are mutants deleted of several normal genes so that the helper viruses and/or their expressed gene products will not cause disease in a  
30 patient.

For example, a helper adenovirus useful in this invention may express only a gene product of a single adenoviral early gene. Exposure of the ss rAAV to an Ad early gene product is sufficient to substantially enhance  
35 the formation of ds rAAV genome with a coordinate

increase in transduction efficiency. The Ad early genes which are useful in producing this effect are E1, E2a, E4 and functional fragments thereof. However, as demonstrated by the examples below, adenovirus

5 substantially enhances recombinant AAV transduction *in vitro* in a way that is dependent on expression of the E1 and E4 genes of adenovirus and is directly proportional to the appearance of ds replicative forms of rAAV.

One example of a helper virus is an adenovirus  
10 deleted of most of its wild-type early genes and which is capable of expressing only its E4 gene or a functional fragment thereof in the target cell. Among such functional fragments is the ORF 6 of the E4 gene. As described below in the examples, experiments in cell  
15 lines indicate that the ORF6 of the adenoviral E4 gene locus is sufficient to significantly enhance rAAV transduction. Selective expression of the E4-ORF6 product of adenovirus accomplishes a increase in transduction efficiency similar to, but somewhat  
20 attenuated, compared to that produced by exposure to the E1 and E4 gene products in combination. That is, the ORF6 product of E4 is sufficient to enhance the augmentation of rAAV transduction; but this effect is amplified substantially by E1 gene products.

25 Thus, more preferably, exposure of the rAAV to both the expressed E1 and E4 gene products produces a substantial enhancement of the above-described rate limiting step. Therefore, another exemplary helper virus may also contain more than one gene which, upon  
30 expression, facilitates the ss to ds conversion. An example of such a helper virus is an adenovirus which expresses both the E1 and E4 genes, or functional fragments thereof. Still other Ad genes may be expressed by the helper virus, provided that the virus is  
35 sufficiently crippled so that it does not cause disease

in the patient contributing the target cells.

Where the agent which facilitates conversion of ss to ds rAAV is a helper virus, the method of the invention comprises co-infecting the target cell with the rAAV and the helper virus. Such co-infection may occur in the context of *ex vivo* therapy, i.e., manipulations performed on cells extracted from the patient, which cells are reinserted into the patient after the method is performed. Alternatively, the patient may be directly co-infected with the two viruses by conventional means. Delivery of the two viruses to the patient may be directed to a specific organ, or to the general circulatory system. Such delivery methods are described in the art for gene therapy of, e.g., cystic fibrosis [see, e.g., US Patent No. 5,240,846].

*B. The Conversion Agent is a Chemical, Drug or Other Entity that can Activate rAAV Transduction*

In another embodiment of the method of this invention, the conversion agent which contacts the cells infected with the rAAV may be selected from the following classes of known compounds or methods: 1) inhibitors of DNA synthesis such as hydroxyurea, hydrogen peroxide, and other direct or indirect inhibitors of DNA polymerase; 2) chemo-therapeutic agents that induce DNA damage, such as cyclophosphamide, alkylating agents, purine analogs, e.g., 6-thioguanine, etc.; 3) drugs that interfere with DNA modifying enzymes, such as inhibitors of topoisomerase, DNA ligase exonucleases and endonucleases; and 4) agents that nonspecifically enhance transcription, such as sodium butyrate, or agents that stabilize cells, such as DMSO. Also, genotoxic agents such as carcinogens may be employed as the conversion agent. Other methods of inducing disruption or damage to DNA may also be useful as agents capable of facilitating ss to ds conversion of rAAV and maybe selected by one of skill in

the art, including physical methods, such as irradiation. These classes of compounds or methods are believed to result in the conversion from ss to ds rAAV.

According to this embodiment of the method of the invention, the rAAV is again produced conventionally, but not co-infected with a helper virus. The ss rAAV is infected into the target cell, and the infected cell is contacted by the agent in an appropriate manner depending on the identity of the agent. These conversion enhancing agents can be employed in ex vivo treatment of the target cells infected by the rAAV by application directly to the cells. Such application can occur substantially simultaneously, or consecutively, with application of the rAAV gene therapy vehicle. For example, the infected target cell may be subjected to one of the above-listed compounds or drugs for a desired time period. The parameters for contacting the infected cells with the agent may readily be determined by one of skill in the art. These parameters will depend upon whether the method is performed ex vivo or in vivo. For example, the number of ex vivo infected cells to be treated will be considered for the dosage, and timing of such treatment.

Similarly, the physical status of the patient can determine the parameters of delivery of the agent to the patient in vivo. The dosage and amount of the damaging agent may therefore be adjusted by one of skill in the art. Where the agents are typical chemotherapeutic drugs approved for use in humans or animals, such enhanced conversion of rAAV may also occur in vivo by the co-administration of the agent, i.e., the chemotherapeutic drug, and the rAAV gene therapy vehicle to the patient. According to this aspect of the invention, the chemotherapeutic drug would be administered only when the rAAV is administered. Appropriate dosages and amounts of chemotherapeutic drugs

and recombinant gene therapy vehicles and means for determining such amounts are within the skill of the art. However, because the effect of the chemotherapeutic drug will enhance the ss to ds conversion of the rAAV and thus  
5 enhance its efficiency of transduction into the target cells, it is anticipated that lower dosages than the conventional dosages of either or both the drug and the rAAV could be effectively administered.

*C. Conversion Agent May be Part of the rAAV.*

10 In still another embodiment of this invention, a novel "second generation" rAAV may be designed to incorporate the conversion agent into the virus, so that both the transgene and the conversion agent are co-expressed in the target cell. Such a novel recombinant  
15 adeno-associated virus comprises the following components:

(a) the DNA of at least a portion of the genome of an adeno-associated virus which portion is capable of transducing at least two selected genes or  
20 functional fragments thereof into a target cell in the absence of cell division; (b) a first selected gene, i.e., the desired transgene, operatively linked to regulatory sequences directing its expression, and (c) a second selected gene, i.e., the "conversion gene"  
25 operatively linked to regulatory sequences capable of directing expression of said second gene. The "conversion gene" upon expression is capable of facilitating the conversion of the ss rAAV to its ds form upon expression. The first and second genes in this rAAV  
30 are flanked by the AAV DNA, preferably the 5' and 3' ITRs. An embodiment of such a second generation rAAV is provided schematically in Fig. 11. Its DNA sequence is provided in SEQ ID NO: 5.

Another embodiment of such a novel rAAV may  
35 include more than one gene which upon expression has the



ability to facilitate conversion of ss to ds rAAV in the target cell. For example, the novel rAAV described above may also contain an additional selected gene operatively linked to regulatory sequences capable of directing its expression, the additional gene and said second "conversion" gene described above being capable of jointly facilitating the conversion of ss rAAV to its ds form upon expression of both the second and additional genes. In this rAAV, all three genes, i.e., the transgene, the second "conversion" gene and the additional gene are flanked by the AAV DNA.

In one desirable embodiment of a novel rAAV<sub>m</sub> the AAV ITRs flank a selected transgene, and a conversion gene, which is the adenovirus E4 gene or a functional fragment thereof (e.g., the ORF6 sequence). In another embodiment, the novel recombinant expresses three genes, the transgene, the adenovirus E4 gene or a functional fragment thereof and the adenovirus E1 gene or a functional fragment thereof. The E4 and E1 gene products expressed in the target cell with the transgene, together act to facilitate conversion of the ss to ds form of rAAV.

In still another embodiment of the novel rAAV and its use, the regulatory sequences directing expression of the conversion gene, e.g., whether it be a single second gene or more than a single additional gene, may include an inducible promoter. Thus, expression of the conversion gene occurs only in the presence of an inducing agent. Many inducible promoters and companion inducing agents, e.g., steroids such as glucocorticoids, are known to the art and may be readily selected for incorporation into the rAAV and methods of this invention by one of skill in the art with resort to this description.

The method of the invention employing such

"second generation" rAAVs which carry at least one "conversion gene" provides for infecting the target cell with this ss rAAV. Where the promoters directing expression of both the transgene and the conversion gene are constitutive, the infected target cell machinery will direct the expression of the transgene product and conversion gene product. Co-expression in the target cell of the transgene and the "conversion gene" facilitates the conversion of ss rAAV to ds rAAV in the cell, and increases the transduction efficiency, and perhaps stable chromosomal integration, without further method steps.

When the second generation rAAV employed in the method contains the "conversion gene(s)" under the control of inducible promoter(s), the method is slightly altered. Following infection of the target cell by the rAAV, the infected target cell is contacted with a suitable inducing agent, which triggers the inducible promoter to "turn on" production of the conversion gene product. When the inducing agent is removed or stopped, the expression of the conversion gene product is "turned off".

As described above, any prior art rAAV containing a transgene for gene therapy may be used in at least one embodiment of the above methods. The sources, selection and assembly of the various components to generate the rAAV, including the novel rAAV described above, are now conventional and readily accessible to one of skill in this art, given the disclosure contained herein. Such methods employ conventional genetic engineering techniques [See, e.g. Sambrook et al, cited above].

The novel rAAV viruses and the methods of this invention provide efficient gene transfer vehicles for somatic gene therapy and are suitable in pharmaceutical

compositions for *ex vivo* applications and *in vivo* use. When rAAV contain a therapeutic gene, e.g., in place of the *LacZ* transgene illustrated in the exemplary rAAV, AV.CMV*LacZ*, by use of the rAAV and the methods described herein, the therapeutic transgene can be delivered to a patient *in vivo* or *ex vivo* to provide for efficient transduction, and possibly stable integration, of the desired gene into the target cell. Thus, these novel rAAV and the methods described herein can be employed to correct genetic deficiencies or defects. The potential of AAV to efficiently integrate its genome into nondividing cells is currently being exploited in the development of gene therapies based on *ex vivo* transduction of hematopoietic stem cells. *In vivo* application of rAAV is primarily being developed for the treatment of CF where purified stocks of virus are instilled into the airway to transduce the terminally differentiated epithelial cells of conducting airway. The methods and compositions described herein can be used with both types of gene therapy. Another condition suitable for such use includes transduction of the low density lipoprotein (LDL) receptor gene into hepatocytes for the treatment of familial hypercholesterolemia. One of skill in the art can generate any number of rAAV which can be used via the above methods for the treatment of these and other disorders.

For *ex vivo* or for *in vivo* therapy, the rAAV may be used to infect the target cells by suspending the virus particles in a biologically compatible solution or pharmaceutically acceptable delivery vehicle. A suitable vehicle includes sterile saline. Other aqueous and non-aqueous isotonic sterile injection solutions and aqueous and non-aqueous sterile suspensions known to be pharmaceutically acceptable carriers and well known to

those of skill in the art may be employed for this purpose.

The rAAV are administered in sufficient amounts to transfect the desired cells and provide sufficient  
5 levels of expression of the selected transgene to provide a therapeutic benefit without undue adverse, or with medically acceptable, physiological effects which can be determined by those skilled in the medical arts. Conventional and pharmaceutically acceptable routes of *in*  
10 *vivo* administration include direct delivery to the target organ, tissue or site, intranasal, intravenous, intramuscular, subcutaneous, intra $\pi$ dermal, oral and other parental routes of administration. Routes of administration may be combined, if desired.

15 Dosages of the rAAV for the infecting step of the method will depend primarily on factors such as the therapeutic environment, i.e., *ex vivo* or *in vivo*; the condition being treated, the selected gene, the age, weight and health of the patient, and may thus vary among  
20 patients. A therapeutically effective dosage of the rAAV for *ex vivo* treatment will be based upon the multiplicity of infection, which is likely to range from between about 1 to about 10 transducing particles/cell. A therapeutically effective human dosage of the rAAV for *in*  
25 *vivo* infection according to the present invention is believed to be in the range of from about 20 to about 50 ml of saline solution containing concentrations of from about  $1 \times 10^7$  to  $1 \times 10^{10}$  transducing viral particles/ml virus. A preferred human dosage is about 20 ml saline  
30 solution at the above concentrations. The dosage will be adjusted to balance the therapeutic benefit against any side effects. The levels of expression of the selected gene can be monitored to determine the selection, adjustment or frequency of dosage administration.

The effective amount of the facilitating agent to be administered is within the skill of the art to determine and will depend upon the identity of the agent. Known dosages of certain of the classes of chemicals and pharmaceuticals described above may be employed in this method to damage the DNA and facilitate ss to ds conversion of the rAAV. Where the agent is a gene expressed by a helper virus, the amounts of infecting virus should be similar to those amounts described above for the rAAV. Of course, where the agent is a gene present in a second generation rAAV, the identical dosages described above for the rAAV will apply.

Several embodiments of the above-described methods of this invention were confirmed in murine models of rAAV mediated gene transfer to both lung and liver. These experiments demonstrated similarly low levels of gene transfer *in vivo* by rAAV, which was increased several orders of magnitude by coinfection with E1 and E4 expressing adenovirus.

In summary, experiments were conducted to demonstrate that adenovirus enhances rAAV transduction in cultured cells. During the production and characterization of a *lacZ* recombinant AAV generated in 293 cells that were coinfectd with an E1 deleted virus, it was observed that purification of rAAV from lysates was associated with substantial loss of *lacZ* transducing activity when assayed on 293 cells. This drop in rAAV activity was particularly evident in the final step where residual contaminating helper adenovirus was removed by heat inactivation. *LacZ* transducing activity was recovered by adding adenovirus back to the purified stock of rAAV. These data provided the first indication that adenovirus could substantially enhance the transduction efficiency of rAAV.

As described in Example 10, a series of complementation groups were generated by mixing different adenovirus early gene mutants with purified *LacZ* rAAV, referred to as AV.CMVLacZ (see Example 2). These defined mixtures of viruses were analyzed for *LacZ* transduction on Hela cells (See Examples 12 and 13). An E1 deletion rAd H5.CBALP and the E4 deletion mutant dl1004 provided no significant increase in AV.CMVLacZ transduction (Fig. 5A). However, partial activity could be achieved with E1 and E4 mutants that carried less severe deletions. Both dl110 (E1B-55kDa deleted) and dl1010 (ORF6 deleted) enhanced transduction to levels that approached those of Ad5, ts125, and dl802 in terms of the number of positive blue cells, but total  $\beta$ -galactosidase activity was substantially lower (Fig. 5A). These results implicate early regions E1 and E4 in the augmentation of rAAV transduction.

The experiments described below also demonstrate that the novel rAAV which incorporates as its conversion gene, an Ad gene, such as E4, can increase transduction efficiency of the rAAV in the absence of a helper virus. As described in more detail below in Example 15 below, 293 cells were stably transfected with a genomic fragment of Ad5 spanning E4. This E1/E4 expressing cell line and the parent E1 expressing cell line (293) were infected with rAAV and analyzed for transduction. These experiments demonstrated the significance of the combined expression of E1 and E4(ORF6) in the adenovirus mediated augmentation of rAAV transduction.

In the presence of E1 and E4 expression, rAAV transduction was invariably accompanied by the appearance of ds RF monomers and dimers (Example 14). Importantly, the tight correlation between rAAV vector transduction and the accumulation of duplex forms could be achieved in

two different experimental settings; cells infected with E1/E4 expressing adenovirus (Figs. 8A and 8B), or complementing cell lines (Fig. 8C).

The following examples illustrate the construction and testing of the novel packaging cell lines, the E1/E4 deleted rAd of the present invention and the use thereof, improved methods and second generation recombinant AAV production for gene therapy of the present invention. These examples are illustrative only, and do not limit the scope of the present invention.

Example 1 - Novel E1a/E1b and E4 Expressing Packaging Cell Lines

*A. Construction of E4 ORF 6 Expressing Plasmids*

The entire E4 region from Ad5 or an ORF6 minigene were subcloned into a shuttle plasmid that contained a neomycin resistance gene. Two versions of ORF6 minigene were developed that differed in the promoter element. The first used a Zn+2 inducible sheep metallothionine (MT) promoter to drive ORF 6 expression. The second used a dexamethasone-inducible mouse mammary tumor virus (MMTV) promoter.

An exemplary plasmid useful for the construction of a packaging cell line of this invention is pMMTVE4ORF6. The minigene contained in this plasmid is set out in SEQ ID NO: 1, and contains a mouse mammary tumor virus promoter (MMTV) (nucleotides 1-1506 of SEQ ID NO:1) in transcriptional control of a human E4 ORF 6 gene sequence (nucleotides 1523-2408 of SEQ ID NO: 1), a growth hormone terminator (GH) (nucleotides 2409-3654 of SEQ ID NO: 1), an SV40 origin of replication, plasmid sequences from plasmid pBR322, including a neomycin resistance gene, and an ampicillin resistance gene. The amino acid sequence of ORF 6 is indicated in SEQ ID NO: 2. The various functional fragments of this plasmid may be readily replaced with other conventionally used

sequences and are not critical to the design of the plasmid.

Another plasmid useful for the construction of a packaging cell line of this invention is pMTE4ORF6.

5 The DNA sequence of the minigene contained in this plasmid is similar to that of SEQ ID NO: 1, except that the promoter is a sheep metallothionine promoter (MT promoter) [M. G. Peterson et al, cited above].

A plasmid used as a control for the  
10 construction of a packaging cell line of this invention is pLTR.E4(-). This plasmid contains the endogenous constitutive retroviral MLV LTR and most of the Ad E4 gene region except that the endogenous E4 promoter and a portion of E4 ORF1 are missing. The other plasmid  
15 sequences remain the same as described above.

Still another plasmid useful for the study of the methods of this invention is pLTR.E4, which contains the constitutive MLV LTR and endogenous E4 promoter and an intact E4 gene. The other plasmid sequences remain  
20 the same as described above.

To determine whether ORF6 expression was sufficient to enhance rAAV transduction, the inducible metallothionein (MT)-ORF6 minigene was stably transfected into HeLa cells. This new cell line, HeLa(MT-ORF6) was  
25 evaluated for LacZ rAAV transduction in response to ORF6 induction as described below. The cell line 293 (MT-ORF6) expresses ORF-6 of the E4 gene of Ad5 from the metallothionine promoter which is relatively inactive at baseline but can be induced with divalent cations. These  
30 293 cells were included to establish the baseline transduction efficiency.

#### B. Transfections and Selection of Clones

Each of the above-described plasmids was transfected by the calcium phosphate precipitation  
35 technique into the human embryonic kidney cell line 293



[ATCC CRL1573] which express the product of the adenovirus E1 genes, or into HeLa cells, seeded on 100 mm plates (10 µg plasmid/plate). Twenty four hours post-transfection, cells were harvested and seeded at varying dilutions (1:10 - 1:100) in 100 mm plates for about 10 days. Seeding media contain G418 (Geneticin, BRL) at 1 mg/ml. Resistant colonies that developed were selected using the following assays and expanded. Preliminary analysis of clones was based on enhanced transduction efficiency of a recombinant adeno-associated virus, AV.CMVLacZ, and immunofluorescence localization of Ad E4 protein as described in the following examples.

Example 2 - Recombinant AAV and AV.CMVLacZ Transduction Enhancement Assay

E1 and E4 Ad gene products are needed for recombinant adeno-associated virus (AAV) function. This primary assay involves seeding the packaging cell lines of Example 1 in 96 well 35 mm culture plates ( $2 \times 10^6$  cells/well) and infecting the cells with purified, heat-treated AV.CMVLacZ at an MOI of 1000 virus particles/cell.

A. Preparation of Recombinant AV.CMVLacZ

A recombinant AAV virus was prepared by conventional genetic engineering techniques for the purposes of this experiment. Recombinant AAV was generated by plasmid transfections in the presence of helper adenovirus [Samulski et al, *J. Virol.*, 63:3822-3828 (1989)]. A cis-acting plasmid pAV.CMVLacZ [SEQ ID NO: 4] (see Fig. 10) was derived from psub201 [Samulski et al, *J. Virol.*, 61:3096-3101 (1987)] and contains an *E. coli* β-galactosidase minigene in place of AAV Rep and Cap genes. The 5' to 3' organization of the recombinant AV.CMVLacZ genome (4.9 kb) [SEQ ID NO: 4] includes

(a) the 5' AAV ITR (bp 1-173) was obtained by PCR using pAV2 [C. A. Laughlin et al, *Gene*,

23: 65-73 (1983)] as template [nuc. 53-219];

(b) a CMV immediate early enhancer/promoter [Boshart et al, Cell, 41:521-530 (1985)] (nuc. 246-839);

5 (c) an SV40 intron (nuc. 856-987);

(d) *E. coli*  $\beta$ -galactosidase cDNA (nuc. 1039-4512);

(e) an SV40 polyadenylation signal (a 237 Bam HI-BclI restriction fragment containing the  
10 cleavage/poly-A signals from both the early and late transcription units (nuc. 4522-4719) and

(f) 3'AAV ITR, obtained from pAV2 as a SnaBI-BglII fragment (nuc. 4759-4925). All other nucleotides are plasmid derived.

15 Rep and Cap genes were provided by a trans-acting plasmid pAAV/Ad [Samulski et al, cited above].

Monolayers of 293 cells grown to 90% confluency in 150 mm culture dishes ( $5 \times 10^7$  cells/plate) were infected with H5.CBALP at an MOI of 10. H5.CBALP (also  
20 called H5.010ALP) is a rAd that contains an alkaline phosphatase minigene in place of adenovirus Ela and Elb gene sequences (map units 1-9.2 of the Ad5 sequence of GenBank [Accession No. M73260]). The alkaline phosphatase cDNA is under the transcriptional control of  
25 a CMV-enhanced  $\beta$ -actin promoter in this virus. This helper virus is described in Goldman et al, Hum. Gene Ther., 6:839-851 (July, 1995); Engelhardt et al, Hum. Gene Ther., 5:1217-1229 (October, 1994); and references cited therein.

30 Infections were done in Dulbecco's Modified Eagles Media (DMEM) supplemented with 2% fetal bovine serum (FBS) at 20 ml media/150 mm plate. Two hours post-infection, 50  $\mu$ g plasmid DNA (37.5  $\mu$ g trans-acting and 12.5  $\mu$ g cis-acting) in 2.5 ml of transfection cocktail  
35 was added to each plate and evenly distributed.

Transfections were calcium phosphate based as described [B. Cullen, Meth. Enzymol., 152:684-704 (1987)]. Cells were left in this condition for 10-14 hours after which the infection/transfection media was replaced with 20 ml fresh DMEM/2% FBS. Forty to fifty hours post-transfection, cells were harvested, suspended in 10 mM Tris-Cl (pH 8.0) buffer (0.5 ml/150 mm plate) and a lysate prepared by sonication. The lysate was brought to 10 mM manganese chloride, after which bovine pancreatic DNase I (20,000 units) and RNase (0.2 mg/ml final concentration) were added, and the reaction incubated at 37°C for 30 minutes. Sodium deoxycholate was added to a final concentration of 1% and incubated at 37°C for an additional 10 minutes.

The treated lysate was chilled on ice for 10 minutes and solid CsCl added to a final density of 1.3 g/ml. The lysate was brought to a final volume of 60 ml with 1.3 g/ml CsCl solution in 10 mM Tris-Cl (pH 8.0) and divided into three equal aliquots. Each 20 ml sample was layered onto a CsCl step gradient composed of two 9.0 ml tiers with densities 1.45 g/ml and 1.60 g/ml.

Centrifugation was performed at 25,000 rpm in a Beckman SW-28 rotor for 24 hours at 4°C. One ml fractions were collected from the bottom of the tube and analyzed on 293 or 293(E4) cells for *LacZ* transduction. Fractions containing peak titers of functional AV.CMV*LacZ* virus were combined and subjected to three sequential rounds of equilibrium sedimentation in CsCl. Rotor selection included a Beckman NVT-90 (80,000 rpm for 4 hours) and SW-41 (35,000 rpm for 20 hours). At equilibrium, AV.CMV*LacZ* appeared as an opalescent band at 1.40-1.41 g/ml CsCl. Densities were calculated from refractive index measurements. Purified vector was exchanged to 20 mM HEPES buffer (pH7.8) containing 150 mM NaCl (HBS) by dialysis and stored frozen at -80°C in the

presence of 10% glycerol or as a liquid stock at -20°C in HBS/40% glycerol.

Purified virus was tested for contaminating H5.CBALP helper virus and AV.CMVLacZ titers. Helper virus was monitored by histochemical staining for reporter alkaline phosphatase activity. A sample of purified virus representing 1.0% of the final product was added to a growing monolayer of 293 cells seeded in a 60 mm plate. Forty-eight hours later, cells were fixed in 0.5% glutaraldehyde/phosphate buffered saline (PBS) for 10 minutes at room temperature, washed in PBS (3x10 minutes) and incubated at 65°C for 40 minutes to inactivate endogenous alkaline phosphatase activity. The monolayer was allowed to cool to room temperature, rinsed once briefly in 100 mM Tris-Cl (pH9.5)/100 mM NaCl/5mM MgCl, and incubated at 37°C for 30 minutes in the same buffer containing 0.33 mg/ml nitroblue tetrazolium chloride (NBT) and 0.165 mg/ml 5-bromo-4-chloro-3-indolylphosphate p-toluidine salt (BCIP). Color development was stopped by washing the monolayer in 10 mM Tris-Cl (pH 8.0)/5 mM EDTA. Routinely the purification scheme described above removed all detectable H5.CBALP helper virus by the third round of buoyant density ultracentrifugation.

AV.CMVLacZ titers were measured according to genome copy number (virus particles/ml), absorbance at 260 nm ( $A_{260}$  particles/ml) and LacZ Forming Units (LFU/ml). Virus particle concentrations were based on Southern blotting. Briefly, a sample of purified AV.CMVLacZ was treated with capsid digestion buffer (50 mM Tris-Cl, pH 8.0/1.0 mM EDTA, pH 8.0/0.5% SDS/Proteinase K 1.0 mg/ml) at 50°C for one hour to release virus DNA. The reactions were allowed to cool to room temperature, loading dye was added and electrophoresed through a 1.2% agarose gel. Standard

quantities of ds AV.CMVLacZ genome were also resolved on the gel.

DNAs were electroblotted onto a nylon membrane, hybridized with a  $^{32}\text{P}$  random primer labeled restriction fragment, and the resulting blot scanned on a PhosphorImager 445 SI (Molecular Dynamics). A standard curve was generated from the duplex forms and used to extrapolate the number of virus genomes in the sample. LFU titers were generated by infecting indicator cells with limiting dilutions of virus sample. Indicator cells included HeLa and 293 and 293 (E4) lines (described in Example 10 below). Twenty-four hours later, cells were fixed in glutaraldehyde and cells were histochemically stained for *E. coli*  $\beta$ -galactosidase (*LacZ*) activity as described in J. M. Wilson et al, Proc. Natl. Acad. Sci. USA, 85:3014-3018 (1988). One LFU is described as the quantity of virus that is sufficient to cause visually detectable  $\beta$ -galactosidase expression in one cell 24 hours post-infection.

#### 20 B. Induction of ORF6 Expression

Induction of ORF6 expression with 10  $\mu\text{M}$  dexamethasone or 150 $\mu\text{M}$  zinc sulfate (for negative control, no inducer used) was initiated 2 hours before the addition of virus and continued throughout the duration of the experiment. Twenty-four hours after the addition of virus, cells were harvested, lysates were generated by sonication and analyzed for the  $\beta$ -galactosidase expression (i.e.,  $\beta$ -galactosidase activity) and virus DNA as described above. Hirt extracts were prepared from low molecular weight DNA from cell extracts. The preparation of the Hirt extracts and subsequent analysis by Southern hybridization were performed by resort to conventional procedures known to on of skill in the art.

35 In the absence of the inducers, the packaging

cell lines generate lower levels of  $\beta$ -galactosidase in rAAV infected cells. Induction of ORF6 expression with the inducer dexamethasone results in a concomitant rise in AV.CMV $\text{LacZ}$  cell transduction to a level that was much greater than the parent 293 line. Expression of E1 alone was insufficient to have an effect in the adenovirus mediated augmentation of rAAV transduction.

Results are demonstrated for certain positive clones in the Table I below (see Example 4). However, for 30 cell lines having an MMTV promoter and ORF6 sequence, 4 demonstrated over 90% blue cells illustrative of LacZ production in the presence of dexamethasone, i.e., 293-27-6, 293-27-17, 293-27-18 and 293-27-28.

Example 3 - Immunofluorescence Localization of Ad5 Late Protein.

Positive clones from the assay of Example 2 were infected with the recombinant E4 deleted adenovirus H5dl1004 and screened for E4 complementation using an immunofluorescence assay for late gene expression. The H5dl1004 virus was obtained from Dr. Ketner of Johns Hopkins University and is described in Bridge and Ketner, J. Virol., 632(2):631-638 (Feb. 1989), incorporated by reference herein. Because ORF6 of E4 complements late Ad gene expression, specifically in the formation of the hexon and penton fibers of the adenovirus, cell lines containing ORF6 are able to bind with antibody against these proteins.

Each cell line of Example 1 is infected with E4 deleted virus H5dl1004 virus at an MOI of 0.1. The cells were treated with mouse anti-adenovirus FITC-labeled monoclonal antibody to either the hexon or penton fibers in a 1:10 dilution (Chemicon International Inc., Temecula, CA). Positive clones were identified by reaction with the antibody.

Example 4 - Relative Plaquing Efficiency

The cell lines of Example 1 demonstrating with strong complementation ability in Example 3 were screened for relative plaquing efficiency of H5dl1004 as compared to W162 cells (an E4-complementing Vero cell line which does not express E1) [Weinberg and Ketner, Proc. Natl. Acad. Sci. USA, 80(17):5383-5386 (1983)]. In Table II below, RPE%, i.e., relative plaquing efficiency, represents the titer of H5dl1004 on tested cell lines/titer of H5dl1004 on W162 cells. For example, the RPE of 293 cells is 0.

The positive cell lines selected by all criteria are identified in Table I below, with the results of the assays of Examples 2, 3 and 4.

15

TABLE I

E1/E4 Double ComPLEMENTING Cell Lines

Cell Line	Trans-Gene	Pro-moter	IF/LP	AV.CMV LacZ	RPE%
293-10-3	ORF6	MT	++++	++++	246
293-39-11	ORF6	LTR	++++	+++	52
293-84-31	E4-	LTR	++++	++++	179
293-12-31	whole E4	LTR +E4	++++	++++	174
293-27-6	ORF6	MMTV		+++++	327
293-27-17	ORF6	MMTV		++++	313
293-27-18	ORF6	MMTV		+++++	339
293-27-28	ORF6	MMTV		++++	261

25

Example 5 - Construction and Purification of H5.001CBLacZ

30

The plasmid pAd.CBLacZ was constructed as described in detail in K. Kozarsky et al, Som. Cell Mol. Genet., 19(5): 449-458 (1993), incorporated by reference herein. This plasmid contained a minigene comprising a 5'

flanking *NheI* restriction site, followed by Ad5 sequence m.u. 0-1, followed by an *E1* deletion into which is inserted a CMV enhancer/chicken  $\beta$ -actin promoter sequence [T. A. Kost et al, Nucl. Acids Res., 11(23):8287 (1983)],  
5 which controls the transcription of the following bacterial  $\beta$ -galactosidase, followed by a poly A sequence and flanked 3' by Ad m.u. 9-16, and another *NheI* site. In the plasmid, the minigene was flanked on both sides by plasmid sequence containing drug resistance markers.

10 The plasmid pAd.CBLacZ was linearized with *NheI* and co-transfected by the calcium phosphate co-transfection method into the novel packaging cell line of Example 1 with *ClaI* digested H5dl1004 (an Ad5 sequence deleted of from about map unit 92.1 through map unit 98,  
15 corresponding to substantially the entire *E4* gene).

Homologous recombination occurs in the cell line between these two viral constructs between Ad map units 9-16, resulting in rAd, designated H5.001CBLacZ [SEQ ID NO: 3] (Fig. 2). This rAd contains the sequence from  
20 pAd.CBLacZ (including Ad map units 0-1 (nuc. 1-330); CMV enhancer/chicken  $\beta$ -actin promoter (CB) (nucs. 370-928); *E. coli*  $\beta$ -galactosidase (nucs. 945-4429); the polyA (nuc. 4429-4628); and Ad5 map units 9-92.1 and 97.3 to 100 from H5dl1004 (nucs. 4671-35408)). This rAd is thereby  
25 functionally deleted, and substantially structurally deleted, of the Ad *E1* and *E4* genes.

Viral plaques were selected and screened by the  $\beta$ -galactosidase assay [Wilson (1988), cited above] and H5.001CBLacZ was isolated following three rounds of  
30 plaque purification. The purified virus was also subjected to cesium chloride density centrifugation and large scale production.

For the following mouse experiments, virus was used after column purification and glycerol was added to a



final concentration of 10% (v/v). Virus was stored at -70°C until use.

Example 6 - Growth Kinetics of H5.001CBLacZ in Packaging Cell Lines

5 The cell lines identified in Table I were infected with recombinant H5.001CBLacZ at an MOI of .5. The growth kinetics of this virus in the E4 complementing cell lines are shown in Fig. 3. Maximum viral yield is reported as LFU/ml in Table II below.

10

TABLE II

	<u>Cell Line</u>	<u>Maximum Viral Yield</u>
	293-10-3	$2.8 \times 10^{10}$
	293-39-11	$9.5 \times 10^8$
	293-84-31	$1.1 \times 10^9$
15	293-12-31	$4.5 \times 10^8$
	293-27-6	$2.8 \times 10^{10}$
	293-27-17	$2.5 \times 10^{10}$
	293-27-18	$2.9 \times 10^{10}$
	293-27-28	$1.2 \times 10^{10}$

20 When grown in 293-27-18 cells (the E4 ORF6 cell line with MMTV promoter inducible by dexamethasone) the maximum yield of this virus is  $2.9 \times 10^{10}$  LFU/ml. Several of the cell lines were passaged between 5 and 20 times and the viral production of the passages remained stable.

25 However, RPE did fall following repeated passages of cells.

Example 7 - Other Recombinant Adenoviruses

Other related rAds were prepared similarly to H5.001CBLacZ by homologous recombination between

30 pAdCBLacZ and other helper viruses.

As one example, H5.000CBLacZ is a recombinant E1 deleted Ad5 which contains the same minigene as H5.001CBLacZ, but has an intact E4 gene. This rAd was prepared as described by homologous recombination between

35 pAdCBLacZ and a wild-type Ad5.

As another example, H5.010CBLacZ contains the adenovirus map units 0-1, followed by a CMV enhanced, chicken cytoplasmic  $\beta$ -actin promoter, the *E. coli*  $\beta$ -galactosidase gene (lacZ), a polyadenylation signal (pA), and adenovirus type 5 map units 9-100, with a small deletion in the E3 gene (the Ad 5 sub360 backbone). This rAd may be prepared by homologous recombination between the pAdCBLacZ vector and Ad5 virus sub360, which contains a 150 bp deletion within the 14.6 kD protein of the E3 gene. See, e.g., J. F. Engelhardt et al, Proc. Natl. Acad. Sci., USA, 91:6196-6200 (June 1994); and Engelhardt et al, Hum. Gene Ther., 5:1217-1229 (Oct. 1994), both incorporated by reference herein.

These rAds were isolated following transfection [Graham, Virology, 52:456-467 (1974)], and were subjected to two rounds of plaque purification. Lysates were purified by cesium chloride density centrifugation as previously described [Englehardt et al, Proc. Natl. Acad. Sci. USA, 88:11192-11196 (1991)]. Cesium chloride was removed by passing the virus over a BioRad DG10 column using phosphate-buffered saline.

#### Example 8 - LacZ Gene Transfer into Mouse

##### A. Transfer into Mouse Muscle

Five to six-week old male C57B/6 mice were anesthetized. Anterior tibialis muscles were exposed and directly injected with either rAd H5.000CBLacZ, H5.010CBLacZ or H5.001CBLacZ as follows: 25  $\mu$ L of purified viral suspension at a stock concentration of  $5 \times 10^{11}$  virus particles/mL was injected by inserting the tip of the 33 gauge needle of a 100  $\mu$ L Hamilton syringe into the belly of the muscle.

Animals were sacrificed on day 4, 14, 28 and 60 post injection. The muscles were dissected and frozen in liquid nitrogen cooled isopentane. Six  $\mu$ M sections were cut in a cryostat, fixed and stained for  $\beta$ -galactosidase

activity for 6 hours at 37°C.

While the blue stained rAd was found for each virus in the day 4 and day 14 (most abundant) stains, by day 28, the H5.001CBLacZ clearly demonstrated more virus on day 28. By day 60, the only virus which stained positive was the H5.001CBLacZ.

B. *Transfer into Mouse Lung and Circulation*

RAd H5.000CBLacZ (control), and H5.001CBLacZ ( $1 \times 10^{11}$  viral particles) were administered to six week old C57BL/6 female mice by tail vein injection and trachea installation. The animals were sacrificed and their liver and lung tissues were harvested at days 4, 9, 21, 28 and 35 post-administration. The transgene and viral late gene expression were compared.

At therapeutic doses of virus, there was diminished expression of late viral proteins at all time points in comparison with transgene.

C. *Dose Responses in Liver*

Dose responses of E4-deleted and E4 intact rAds in the liver of C57BL/6 mice were studied by tail vein administration of  $1.5 \times 10^{11}$ ,  $5 \times 10^{10}$ ,  $1.7 \times 10^{10}$ ,  $5.6 \times 10^9$ , and  $1.9 \times 10^9$  viral particles and comparing the transgene and viral late gene expression at day 4, 21, 28, 35, and 42 post administration.

At therapeutic doses of virus, there was diminished expression of late viral proteins at all time points in comparison with transgene.

Example 9 - Other Gene Transfers

A. *Human OTC Gene Transfer*

The human OTC gene [A. L. Horwich et al, *Science*, 224:1068-174 (1984)] or the human CFTR gene [Riordan et al, *Science*, 245:1066-1073 (1989)] was used to replace the *LacZ* as the transgene in the recombinant E1/E4 deleted adenoviruses described above, using the techniques analogous for the construction of the above-

described *LacZ* vectors.

The resulting human OTC-containing rAd were administered at an MOI of 10 to 30 to human hepatocytes. The E1/E4 deleted rAd demonstrated less replication and less late gene expression than when the E1/E4 deleted rAds are administered to muscle, as described in the example above. However, the results of this gene transfer are better than comparable transfers with rAds containing only a deletion in the E1 gene or a deletion in the E1 gene and a point mutation in the E2a gene.

Similar results are demonstrated when the transgene is CFTR and the method of administration is intratracheal into lungs.

Example 10 - Transduction Efficiency of rAAV *LacZ* AV.CMV*LacZ*) in HeLa Cells Infected with Ad Mutants

A. Viruses

The following viruses were employed in this experiment:

- (1) Wild-type Ad 5, propagated in 293 cells;
- (2) Ad dl110 (an Ad which is deleted of the 55 kb E1B gene) [Babiss et al, J. Virol., 52(2):389-395 (1984) and Babiss and Ginsberg, J. Virol., 50(1):202-212 (1984)], propagated in 293 cells,
- (3) H5.CBALP (an Ad deleted of its E1A and E1B genes and containing a minigene that expresses alkaline phosphatase from a CMV enhanced  $\beta$ -actin promoter, as described above), propagated in 293 cells,
- (4) Ad ts125 (an Ad with a temperature sensitive mutation in the E2A gene which encodes the DNA binding protein) [Ensinger and Ginsberg, J. Virol., 10(3):328-339 (1972)], propagated in 293 cells,
- (5) Ad dl802 (an Ad deleted of its E2a gene), grown in E2A-complementing gmDBP cells as described in Rice and Klessig, J. Virol., 56(3):767-778 (1985);

(6) Ad dl1004 (an Ad deleted of the E4 gene), grown in E4-complementing Vero W162 cells [Weinberg and Ketner, Proc. Natl. Acad. Sci. USA, 80(17):5383-5386 (1983)] and

- 5 (7) Ad dl1010 (an Ad deleted of ORF6 of its E4 gene), grown in E4-complementing Vero W162 cells [Weinberg and Ketner, cited above].

All viruses were purified by two sequential rounds of buoyant density ultracentrifugation in CsCl.

10 B. *Experimental Procedures*

HeLa cells seeded in 6 well, 36mm culture plates ( $2 \times 10^6$  cells/well) were infected with wild-type Ad5 or an adenovirus early gene mutant as described in Part A at an MOI of 10pfu/well. Infections were done in 15 1.0 ml DMEM/2%FBS. Six hours post-infection, monolayers were washed and 1.0 ml fresh DMEM/2% FBS media containing AV.CMVLacZ at  $4 \times 10^9$  virus particles/ml were added. Although the AV.CMVLacZ virus lot used in these experiments was shown to be free of H5.CLALP helper virus 20 by histochemical staining, the virus sample was subjected to heat treatment (60°C for 20 minutes) prior to use to ensure the absence of contaminating adenovirus. Two hours later, 1.0 ml of DMEM/115% FBS was added to each well.

25 Twenty-four hours after the addition of AV.CMVLacZ, cells were harvested. Each test condition was done in triplicate to enable virus transduction to be evaluated in terms of three outputs: histochemical staining for  $\beta$ -galactosidase activity (below), 30 intracellular  $\beta$ -galactosidase specific activity (Example 11), and the molecular form of the virus DNA (Example 12).

HeLa cells were histochemically stained for *E. coli*  $\beta$ -galactosidase (LacZ) activity as described in J. 35 M. Wilson et al, Proc. Natl. Acad. Sci. USA, 85:3014-3018

(1988). The different combinations that were tested included cells transfected with AAV vector alone (AV.CMVLacZ), vector plus wild-type Ad5 (+Ad5), vector plus dl110 (+dl110), vector plus Ad mutant H5.CBALP (+H5.CBALP), vector plus Ad mutant ts125 (+ts125), vector plus Ad mutant dl802 (+dl802), vector plus Ad mutant dl1004 (+dl1004), and vector plus Ad mutant dl1010 (+dl1010).

The results were observed in photomicrographs at magnification 10X (not pictured) of histochemical stains for recombinant  $\beta$ -galactosidase activity. The results indicated that wild-type Ad5 and the E2a mutants ts125 and dl802 caused a significant increase in LacZ rAAV transduction as measured by the number of positive blue cells and the degree of stain intensity. Both dl110 (E1B-55kDa) and dl1010 (ORF6) enhanced transduction to levels that approached those of Ad5, ts125, and dl802 in terms of the number of positive blue cells.

The E1 deletion recombinant H5.CBALP provided no significant increase in AV.CMVLacZ transduction. Expression of E1 alone was insufficient to have an effect in the adenovirus mediated augmentation of rAAV transduction as evidenced by lack of significant increase in transduction obtained with HeLa cells infected with the E4 deletion mutant dl1004. A significant drop in transduction occurred following removal of ORF6 from the E4 region from the coinfecting adenovirus (Fig. 5A).

It is believed that these results demonstrate that the adenoviral gene products, E4 and E1 indirectly promote the formation of ds DNA intermediates that are transcriptionally active.

#### Example 11 - Quantitation of Enhanced Vector Transduction

(A) A duplicate set of HeLa cells as described in Example 10B were used in this experiment. Twenty-four hours after the addition of AV.CMVLacZ recombinant, for

intracellular  $\beta$ -galactosidase assays, cell pellets were suspended in 0.5 ml PBS and sonicated. Cell debris was removed by centrifugation (15,000Xg for 10 minutes) and the clarified extract assayed for total protein [M.

5 Bradford, Anal. Biochem., 72(1-2):248-254 (1976) and M. Bradford et al, Fed. Proc., 35(3):274 (1976)] and  $\beta$ -galactosidase activity [Sambrook et al, cited above] using o-nitrophenyl  $\beta$ -D-galactopyranoside (ONPG) as substrate.

10 Fig. 5A demonstrates the transduction efficiency quantitated by measuring  $\beta$ -galactosidase enzyme activity in the lysates from infected Hela cells and also assayed for total protein. In Fig. 5A, the test condition is shown along the horizontal axis, and  
15 intracellular  $\beta$ -galactosidase specific activity (milliunits/mg protein) using ONPG as substrate is plotted on the vertical axis. Below each bar, the fold-induction in specific activity relative to cells that received the AV.CMVLacZ vector alone is given.

20 The results of Fig. 5A demonstrate that the E2a mutants ts125 and dl802 produced 134-fold and 225-fold increases in  $\beta$ -galactosidase activity, respectively, as compared to that achieved with purified rAAV alone. In comparison, cells infected with wt Ad5 generated 107-fold  
25 increase in  $\beta$ -galactosidase activity.

(B) In another experiment, HeLa cells ( $2 \times 10^6$ ) were infected with increasing multiplicities of wild-type Ad5 or the E2 mutant dl802. Six hours post-infection, monolayers were washed and infected with AV.CMVLacZ at  
30 1000 virus particles/cell. Twenty-four hours after the addition of AV.CMVLacZ, cells were harvested and assayed for total protein and  $\beta$ -galactosidase activity.

The results are illustrated in the bar graph of Fig. 5B, in which adenovirus MOI's are given along the  
35 horizontal axis, and intracellular  $\beta$ -galactosidase

specific activity along the vertical axis. Enhancement of rAAV transduction was proportional to input helper adenovirus from MOIs of 1 to 50 for both wild type Ad5 and dl802. Higher doses of virus were cytopathic, leading to a fall in  $\beta$ -galactosidase expression. Enhanced transduction was achieved when the cells were infected prior to, or at the time of, rAAV infection. The E1 deletion recombinant H5.CBALP and the E4 deletion mutant dl1004 provided no significant increase in AV.CMV $\text{LacZ}$  transduction. Both cells infected with dl110 (E1B-55kDa) and with dl1010 (ORF6) demonstrated substantially lower total  $\beta$ -galactosidase activity than those infected with Ad5, ts125, or dl802.

Example 12 - Analysis of Low Molecular Weight DNAs in AV.CMV $\text{LacZ}$  Transduced Cells

Studies with these early gene mutants of adenovirus suggested that expression of adenoviral genes rather than the virion itself was responsible for enhancement of rAAV transduction. To further investigate these mechanisms and to determine if conversion of ss to ds genome limits the transduction efficiency of rAAV, the molecular state of the rAAV genome was characterized in the infected cells. The relationship between RFm formation and  $\text{lacZ}$  rAAV transduction was explored in experiments where the dose of coinfecting virus was varied (MOI=1, 5, or 10).

(A) A duplicate set of HeLa monolayers as described in Example 10 were harvested 24 hours after they were transduced with the recombinant AV.CMV $\text{LacZ}$  and cultured with or without helper adenovirus.

Episomal DNA was extracted from cell pellets using a modification of the procedure originally described by B. Hirt, J. Mol. Biol., 26:365-369 (1967). Briefly, cells were suspended in 320 ml Tris-Cl (pH8.0)/10 mM EDTA and SDS added to a final concentration of 1%. The mixture was incubated at 37°C for 30 minutes.



Pronase and proteinase K were added to final concentrations of 500 µg/ml and 20 µg/ml, respectively, and the reaction incubated at 37°C for 2 hours. Sodium chloride was added to a final concentration of 1.1 M and  
5 incubated at 4°C overnight. The precipitate that developed during the 4°C incubation was pelleted at 20,000 xg for 30 minutes and the clear supernatant carefully removed. The supernatant was extracted once with phenol:chloroform:isoamyl alcohol (25:24:1) followed  
10 by chloroform:isoamyl alcohol (24:1). Nucleic acids were precipitated with ethanol. The final pellet was suspended in 50 µl Tris-Cl (pH 8.0)/1.0 mM EDTA.

These Hirt extracts were analyzed by Southern blot hybridization. Samples (5 µl) of each Hirt extract  
15 were resolved through a 1.2% agarose gel, electroblotted onto a nylon membrane and hybridized with a <sup>32</sup>P random primer labeled cDNA of the SV40 polyA signal used in AV.CMV<sub>LacZ</sub>.

An autoradiogram of the experiment of Example  
20 12 (not pictured), identifies and labels bands corresponding to the ss AV.CMV<sub>LacZ</sub> genome (SS), a monomer replicative form (RFm), and concatomer replicative forms (RFd). Bands corresponding to the ss AV.CMV<sub>LacZ</sub> genome (SS), a monomer replicative form (RFm), and concatomer  
25 replicative forms (RFd) were identified and labeled. To reference the RFm band, a plasmid carrying AV.CMV<sub>LacZ</sub> was digested to release the entire genome. Autoradiogram exposure times were 14 hours and 69 hours.

In this autoradiogram, the full spectrum of  
30 molecular species present during a lytic infection was demonstrated in cells infected with both *LacZ* rAAV and wild type adenovirus. Both the input ss genome (SS) and monomeric and dimeric forms of ds replicative intermediates (RFm and RFd) are present. This contrasts  
35 with cells infected with purified rAAV alone, where ss

genome is the sole molecular form detected. Analysis of cells coinfecting with the adenovirus early gene mutants revealed a direct correlation between formation of ds forms of the rAAV genome and the enhancement of *LacZ* transduction. Mutant adenoviruses that were ineffective in enhancing rAAV transduction (i.e., the E1 deleted mutant H5.CBALP and the E4 deleted mutant dl1004) failed to promote the formation of ds forms of AAV.

Cells infected with adenovirus deleted of E2a (dl802) or partially deleted of E1 (dl110) or E4 (dl1010) additionally demonstrated a band whose size was identical to the ds replicative monomer (RFm) of the *lacZ* rAAV genome and whose abundance correlated directly with the expression of  $\beta$ -galactosidase activity (compare results of Example 14 to these described results). Slower migrating concatomers, likely dimers, of duplex rAAV were also detected in the autoradiogram described above.

In the presence of E1 and E4 expression, rAd transduction was invariably accompanied by the appearance of ds RF monomers and dimers.

The high molecular weight band in sample lane +H5.CBALP is helper virus DNA. Helper virus DNA is recognized by the SV40 probe because the CBALP minigene also utilizes the SV40 polyA signal.

(B) In another experiment, HeLa cells were infected with wt Ad5 or the E2 deleted mutant dl802 as described in Example 10B. Monolayers were harvested 24 hours later and analyzed for  $\beta$ -galactosidase activity and RFm synthesis. Monomer bands similar to those shown in the autoradiogram described above were quantitated on a PhosphorImager 445 SI and assigned values (CPM).

The results are illustrated in the graphs of Figs. 8A and 8B, in which  $\beta$ -galactosidase specific activity and CPM are plotted along the vertical axis of each figure. Adenovirus MOI's are given on the horizontal

axis of each figure. Data obtained from low MOI infections (1, 5, and 10) are shown. Importantly, the tight correlation between rAAV vector transduction and the accumulation of duplex forms could be achieved in cells infected with E1/E4 expressing adenovirus. The level of  $\beta$ -galactosidase and abundance of RFm increased in proportion to the amount of infecting wild type Ad (Fig. 6A) and dl802 (Fig 6B). These data suggest that synthesis of an episomal duplex intermediate is an obligatory event in transduction.

Example 13 - Duplex End-Analysis

The following is a description of a model for leading strand synthesis of a complementary AAV strand in the presence of Rep (+Rep) or absence of Rep (-Rep). Refer to Figs. 7A-7F. Rep expresses a terminal resolution activity that can convert a duplex structure with closed-ends to an open-ended duplex. In the absence of Rep, terminal resolution is impaired leaving the covalently closed, hairpin structures intact. Under these conditions, hairpins are expected to be found leftward and rightward, since both strands of a rescued ds AAV genome are packaged into virions. Figs. 7B-7F are a flow chart demonstrating the strategy for identifying the terminal structure of duplex RFm that is synthesized from ss AV.CMVlacZ in response to adenoviral gene expression.

Fig. 7C illustrates a closed end and an open end fragment of rAV.CMVlacZ. Figs. 7D, 7E and 7F indicate the mixture of open-ended and covalently closed duplex fragments generated by NotI digestion at position 4509 in the absence of terminal resolution. The NotI 4509 digestion provides a convenient means of releasing a 361 bp fragment that contains the right ITR in the context of a hybridization target (i.e. SV40 pA). In the presence of terminal resolution, only the open-ended 361 bp

fragment would be expected to be generated (Fig. 7D) by such digestion.

The resulting electrophoretic gel (not pictured), revealed in lane (1) the results of digestion of a plasmid carrying an AV.CMVLacZ cDNA to release the rAAV vector, and subsequent digestion with NotI to release the right terminal 361 bp fragment. In lane (2) a sample of NotI digested Hirt DNA extracted from HeLa cells infected with wild-type Ad5 and transduced with AV.CMVLacZ resulted in the release of two fragments, labeled FormI and FormII. (See, also, Figs. 8A and 8B). The migration of ss AV.CMVLacZ (SS) and RFm were also seen.

The ds AV.CMVLacZ intermediates that accumulated in cells infected with adenovirus were likely the result of leading strand DNA synthesis, initiating from the duplex region of the vector ITR. In the absence of Rep, this conversion event was anticipated to generate molecules in which one end is open and the other is covalently closed (Fig. 7A). To further characterize the structure of this ds intermediate Hirt extracts from cells coinfectd with rAV.CMVLacZ and Ad5 were digested with NotI to release the termini of the ds intermediate which, if left open, would be approximately 361 bp in length. The resulting filters were hybridized with a probe specific for the SV40 polyadenylation signal positioned immediately upstream of the rightward ITR. At least two forms were released from the right end of duplex genomes, one that migrated to a position in the gel that predicted an open-ended conformation (Form II), and a second slower migrating species (Form I). Although this result was consistent with the model (Figs. 7A-7F), it was difficult to predict with certainty the structure of Form I. Its retarded mobility did, however, suggest a conformation that differed from the open-ended Form II.

Example 14 - Analysis of AV.CMVLacZ Transduction  
Efficiency in 293 Cells Stably Transfected with an  
Inducible E4 ORF6 cDNA

Cell lines used in this assay were prepared as  
described in Example 1. 293(MT-ORF6) cells and HeLa(MT-  
ORF6) cells were seeded in 6 well 35 mm culture plates  
( $2 \times 10^6$  cells/well) and infected with purified, heat-  
treated AV.CMVLacZ at an MOI of 1000 virus  
particles/cell. Induction of ORF6 expression with from  
none to increasing concentrations of zinc sulfate was  
initiated 2 hours before the addition of virus and  
continued throughout the duration of the experiment.

Twenty-four hours after the addition of virus, cells  
were harvested, lysates were generated by sonication and  
analyzed for the  $\beta$ -galactosidase expression (i.e.,  
 $\beta$ -galactosidase activity) and virus DNA as described in  
the preceding examples. Hirt extracts were prepared from  
low molecular weight DNA from cell extracts. The  
preparation of the Hirt extracts and subsequent analysis  
by Southern hybridization were performed similarly to  
those described in the examples above.

The results of this experiment were as follows:

(1) Specific Activity

The results are illustrated in the bar  
graph of Fig. 8A. Specific activity (milliunits  
 $\beta$ -galactosidase/mg protein) is plotted along the vertical  
axis. Below each bar is given the concentration of zinc  
used for induction, the fold-induction relative to 293  
cells, and the fold-induction relative to 293(ORF6) cells  
maintained in the absence of zinc. As shown in Fig. 8A,  
in the absence of  $Zn^{+2}$ , the 293(MT-ORF6) cell line  
generated 39-fold higher levels of  $\beta$ -galactosidase in  
rAAV infected 293 cells. Induction of ORF6 expression  
with increasing amounts of  $Zn^{+2}$  resulted in a concomitant  
rise in AV.CMVLacZ cell transduction to a level that was

445-fold greater than the parent 293 line. Expression of E1 alone was insufficient to have an effect in the adenovirus mediated augmentation of rAAV transduction.

The specific activity of  $\beta$ -galactosidase was 196.2 mUnits/mg in E1/E4 expressing 293 cells, compared to 1.0 mUnit/mg in 293 cells that only expressed E1 genes. These experiments support a mechanism for enhancing rAAV transduction that is dependent on the combined expression of both E1 and E4 adenoviral genes.

(2) Molecular Analysis of the AV.CMVLacZ Genome

The duplex monomer replicative form (RFm) was quantitated and the values (CPM) plotted along the vertical axis in the bar graph of Fig. 8B. The concentration of zinc used for induction and the fold-induction relative to 293(ORF6) cells maintained in 0 mM zinc is given below each bar.

An autoradiogram (not pictured) shows the agarose gel resolved Hirt extracts from the AV.CMVLacZ transduced cells described above. A plasmid carrying the AV.CMVLacZ cDNA was digested to release the entire sequence and loaded in a lane of the autoradiogram. The band that appeared in this lane therefore reflected the migration of a monomer duplex replicative form (RFm). The migration of the ss AV.CMVLacZ genome (SS), RFm, and dimers of the duplex replicative form (RFd) were also shown. Lanes of the autoradiogram labeled (0), (50), (100), (150), (200), and (250) contained samples from 293(MT-ORF6) cells that were induced with the indicated concentration of zinc. A Hirt extract from 293 cells (lane labeled 293) transduced with AV.CMVLacZ was also shown.

Analysis of Hirt extracts revealed the presence of the RFm in the rAAV infected 293(MT-ORF6) cells that was not present in similarly infected 293

cells. When the induction profiles (Figs. 8A and 8B) that describe AV.CMV $\text{LacZ}$  transduction efficiency were compared, the results were plotted in Fig. 8C. Specific activity (milliunits  $\beta$ -galactosidase/mg protein) data from Fig. 8A and counts-per-minute data (CPM) of AV.CMV $\text{LacZ}$  RfM from Fig. 8B are plotted along the vertical axis, and concentration of zinc sulfate used during the experiment is shown along the horizontal axis.

The two profiles are near mirror images.

Importantly, the RfM increased in proportion to the increment in  $\text{lacZ}$  transducing activity that occurred as ORF-6 expression was induced with  $\text{Zn}^{+2}$  (Fig. 8C). Similar results were obtained with a 293 derived cell line that expresses ORF6 from the glucocorticoid responsive MMTV promoter.

Example 15 - Enhanced AV.CMV $\text{LacZ}$  Transduction in HeLa Cells Carrying an Inducible ORF6 Minigene

HeLa(MT-ORF6) cells ( $2 \times 10^6$ ) were transduced at an MOI of 1,000 AV.CMV $\text{LacZ}$  recombinant particles/cell in absence of zinc sulfate inducer or in the presence of 50, 100, 150, 200, or 250  $\mu\text{M}$  zinc sulfate inducer in the media during transduction. Twenty-four hours later, cells were harvested, cell extracts were prepared by sonication, and analyzed for transgene expression (i.e.,  $\beta$ -galactosidase activity). Cell monolayers were histochemically stained for  $\beta$ -galactosidase activity.

The resulting photomicrographs (not pictured) illustrated that histochemical staining revealed an increase in the number of cells scored  $\text{lacZ}$  positive as the concentration of  $\text{Zn}^{+2}$  in the medium was raised from 0 to 200 mM. Concentrations of 250 mM zinc were found to be toxic to the cells.

Specific activity (milliunits  $\beta$ -galactosidase/mg protein) is plotted in Fig. 9 along the vertical axis.

Below each bar is given the concentration of zinc used

for induction, the fold-induction relative to HeLa cells, and the fold-induction relative to HeLa(MT-ORF6) cells maintained in the absence of zinc. Histochemical staining revealed an increase in the amount of

5  $\beta$ -galactosidase in lysates as the concentration of  $Zn^{+2}$  in the medium was raised from 0 to 200 mM.

Example 16 - Southern Blot Analysis of Low Molecular Weight DNAs from AV.CMV $LacZ$  Transduced HeLa(MT-ORF6) Cells

10 Following Induction of E4ORF6

Hirt extracts were prepared from HeLa(MT-ORF6) cells transduced with AV.CMV $LacZ$  as described in Example 15 in the presence of increasing concentrations of  $Zn^{+2}$  to determine whether synthesis of duplex intermediates

15 contributed to the augmentation in AV.CMV $LacZ$  transduction.

Samples of HeLa(MT-ORF6) cells that were induced with a concentration of zinc sulfate (0, 50, 100, 150, 200, and 250) were resolved on a 1.2% agarose Southern

20 gel (not pictured), transferred to a nylon membrane, and hybridized with a  $LacZ$ -specific probe. One lane contained a plasmid encoding AV.CMV $LacZ$  that was digested to release the entire genome. Bands corresponding to the ss AV.CMV $LacZ$  genome (SS), duplex monomers (RFm), and

25 duplex dimers (RFd) were indicated on the gel.

Southern analysis indicated that HeLa and uninduced HeLa(MT-ORF6) cells demonstrated a single band on Southern blots which comigrated with the ss genome. Induction of ORF-6 resulted in the appearance of

30 detectable levels of ds monomer but only at higher concentrations of  $Zn^{+2}$ . A band comigrating with the RFd was present in all cell preparations, the relevance of which is unclear since the monomer is a likely precursor to the dimer.



Example 17 - Effect of Adenovirus Infection on In Vivo  
AV.CMVLacZ Targeting Efficiency To Murine Liver

The impact of adenoviral gene expression on rAAV transduction in murine liver was studied by sequentially  
5 infusing into the portal vein early gene mutants of adenovirus followed by rAAV.

Balb/c mice, 4- to 6-weeks old [Jackson Laboratories, Bar Harbor, Maine] were anesthetized by an intraperitoneal injection of ketamine (70 mg/kg) and  
10 xylazine (10 mg/kg). For liver studies, a 1 cm left flank incision was made and the spleen exposed.

Samples of purified, heat-treated AV.CMVLacZ in 50  $\mu$ l HBS ( $1 \times 10^{11}$  virus particles) were used alone or spiked with helper adenovirus containing  $2 \times 10^{10}$   $A_{260}$  particles of  
15 purified dl1004, H5.CBALP, or ts125 in a final volume of 50  $\mu$ l. The dose of adenovirus was sufficient to transduce >25% of hepatocytes. The virus mixture was injected just beneath the splenic capsule and the abdomen was closed with 3-0 vicryl.

20 Necropsies were performed 3 days post-infusion and tissue frozen in O.C.T. embedding compound. Frozen sections (6 $\mu$ m) (LacZ+ALP) were prepared and histochemically stained for  $\beta$ -galactosidase enzyme and alkaline phosphatase activity. Sections were  
25 counterstained with neutral red and mounted.

A  $\beta$ -galactosidase positive hepatocyte targeted with AV.CMVLacZ at magnification 20X was obtained. Histochemical analyses of liver tissue harvested 3 days after gene transfer demonstrated that administration of  
30  $10^{11}$  particles of purified rAV.CMVLacZ alone into the portal vein was not associated with appreciable gene transfer (<0.01% of cells), confirming the inherent inefficiency of the rAAV system.

Preinfusion with E4 deleted virus had no impact on  
35 rAAV transduction in mouse liver, whereas E1 deleted

virus demonstrated a modest increment in *lacZ* positive hepatocytes to about 0.1%. The most significant increase in rAAV transduction occurred following infusion of the E2a adenovirus mutant ts125 with *lacZ* expression detected in 10-25% of hepatocytes. A direct relationship between adenovirus gene expression and rAAV transduction was demonstrated in animals infused with both *lacZ* rAAV and the ALP expressing E1 deleted virus. The dose of adenovirus was reduced 10-fold to minimize the coincidental occurrence of coinfection. Histochemical studies demonstrated co-localization of ALP and  $\beta$ -galactosidase in the majority of  $\beta$ -galactosidase expressing hepatocytes.

Example 18 - Effect of Adenovirus Infection on In Vivo AV.CMVLacZ Targeting Efficiency To Murine Lung

Experiments described in Example 17 for mouse liver were adapted for the study of rAAV mediated gene transfer to mouse lung. For lung experiments, anesthetized Balb/C animals were intubated as described in DeMatteo et al, Transplantation (Baltimore), 59(5):787-789 (1995). Briefly, a midline 2 cm skin incision was made in the neck to expose the trachea. A 2 inch 18 gauge angiocatheter was passed through the mouth, positioned in the midportion of the trachea, and connected to a rodent ventilator (#55-3438 Harvard). Polyethylene (PE#10, Intramedic) was fed through the catheter via a side port and advanced beyond the tracheal bifurcation. Using a Hamilton syringe, virus samples (30 $\mu$ l) were slowly infused into the lung through the polyethylene tubing. Samples contained the same formulation of purified, heat-treated AV.CMVLacZ with or without helper adenovirus, as described for liver injections.

Tissue was harvested 72 hours post-infusion. Frozen sections were histochemically stained for  $\beta$ -galactosidase activity and counterstained with neutral red.

Frozen sections from lung (AV.CMVLacZ) showed a  $\beta$ -galactosidase positive airway epithelial cell targeted with AV.CMVLacZ. Similar studies were performed in the murine model of lung-directed gene transfer.

- 5 Adenoviruses were instilled into the trachea prior to the instillation of rAAV. Analysis of lung tissue 3 days later revealed only a rare  $\beta$ -galactosidase positive cell in animals instilled with rAAV alone. No detectable enhancement of rAAV transduction was noted in animals  
10 preinstilled with adenovirus deleted of either E1 or E4. Substantial enhancement of transduction was achieved in conducting airway and alveolar cells of animals administered the E2a mutant adenovirus.

- These experiments in murine models of gene therapy  
15 directed to liver and lung verified that the efficiency of rAAV transduction is low due limited conversion of the input ss genome to a transcriptionally active ds intermediate, and that this conversion is facilitated by expression of adenovirus E1 and E4 gene products.

20 Example 19 - Second Generation rAAV with Regulated Minigene Capable of Enhancing Transduction

- The experiments described in previous examples illustrated the following principles: 1) purified rAAV is a relatively inefficient gene transfer vehicle *in vitro*  
25 and *in vivo* and 2) the rate limiting step in transduction is not viral entry but rather conversion of the virion's ss DNA genome to a transcriptionally active ds DNA genome. Adenovirus can substantially enhance transduction through expression of a subset of its genes.  
30 It does this by promoting conversion of the virion's genome to its ds form. One approach to accomplish this is to incorporate into the recombinant AAV genome a minigene that expresses the minimal adenoviral genes necessary to enhance transduction, i.e., the ORF6 region  
35 f E4.

Two approaches have been considered in designing this modified rAAV. The first strategy is based on a rAAV genome that has two transcriptional units in series, one expressing the therapeutic gene and the other  
5 expressing its E4 ORF6 from a constitutive promoter. While this may, in fact, be useful in many situations, constitutive expression of ORF6 may be detrimental to the cell and potentially could elicit a destructive immune response.

10 The second version of this rAAV includes the therapeutic minigene in addition to the ORF6 transcriptional unit which, in this case, is expressed from an inducible promoter. When this second gene rAAV is administered to the cells (*ex vivo* strategies) or to  
15 the patient (*in vivo* strategies), the inducing agents are administered at the time of gene transfer or soon thereafter. If the ds genomic form or its integrated derivative is stable, the induction of ORF6 will only be necessary at the time of gene transfer into the recipient  
20 cell. Following this, its inducing agent will be withdrawn and the ORF6 gene will be turned off.

An rAAV that illustrates this concept of inducible ORF6 has been constructed and tested *in vitro*. A schematic of the vector pAV.CMVALP.GRE-ORF6, is shown in  
25 Fig. 11 and its sequence is illustrated in SEQ ID NO: 5. This second generation construct contains flanking 5' and 3' AAV ITR sequences. The human placental alkaline phosphatase cDNA (ALP) is included in a minigene in which the promoter from the immediate early gene of  
30 cytomegalovirus drives the transcription. A second transcriptional unit is cloned between the ITRs in series and in direct orientation with the alkaline phosphatase minigene. The second transcriptional unit expresses the Ad5-E4-ORF6 from a glucocorticoid dependent promoter  
35 (GRE) with an SV40 polyadenylation signal. This is

called a second generation rAAV construct.

Specifically, pAV.CMVALP.GRE-ORF6 [SEQ ID NO: 5] generates a novel rAAV containing the *LacZ* transgene and the Ad E4 ORF 6 which facilitates ss to ds conversion of rAAV. The plasmid includes a flanking AAV 5' ITR sequence (nucs. 53-219); CMV enhancer/promoter (nucs. 255-848); human placenta alkaline phosphatase cDNA (ALP) (nucs. 914-2892); SV40 polyA (nucs. 2893-3090); GRE promoter (nucs. 3114-3393); Ad5 E4-ORF6 cDNA (nucs. 3402-4286); SV40 polyA (nucs. 4315-4512); and 3' AAV ITR (nucs. 4547-4713). All other nucleotides are plasmid derived.

The second generation rAAV construct was used to produce and purify rAAV virions which were exposed to HeLa cells that were left untreated or incubated with dexamethasone. In the absence of dexamethasone, (a condition under which little ORF6 should be expressed), little transduction was observed as measured by expression of the alkaline phosphatase gene. Cells incubated in dexamethasone expressed in ORF6 gene and the transduction efficacy was enhanced at least 5-fold. This provides evidence to support that a gene product expressed from the rAAV can function in cis to enhance expression of the transgene.

#### Example 20 - Application to Bone Marrow Directed Gene Therapy

Bone marrow directed gene therapy represents the paradigm of ex vivo gene therapy where the target cell is the hematopoietic stem cell. The basic strategy is to incorporate (i.e., integrate) a therapeutic minigene into the chromosomal DNA of hematopoietic stem cells which are transplanted into a recipient patient whose own bone marrow has been ablated allowing repopulation of its lymphohematopoietic system with progeny of the genetically corrected stem cell.

The problem with this approach has been efficiently transfecting genes into stem cells. Most studies of bone marrow directed gene therapy have utilized recombinant retroviruses which have not been very efficient. One  
5 problem is that retroviruses integrate their provirus only when the target cell is dividing. Unfortunately, most stem cells *in vitro* are quiescent and not dividing. rAAV holds the promise of integrating the provirus more efficiently into non-dividing stem cells. However,  
10 purified rAAV is not very efficient with respect to integration when used alone. In cultured cells, integration is observed in less than 1% of the cells. The same conditions that activate the conversion of ss to ds genome also enhance the integration of the ds  
15 intermediate into the chromosomal DNA.

Therefore, a desirable application of the methods and compositions of this invention is in bone marrow directed gene therapy. According to this method, stem cells are genetically modified with rAAV and an inducing  
20 agent *ex vivo* using the constructs and methods described above (see e.g., Example 19). Genetically modified stem cells are subsequently transplanted by conventional techniques.

Numerous modifications and variations of the present  
25 invention are included in the above-identified specification and are expected to be obvious to one of skill in the art. Such modifications and alterations to the compositions and processes of the present invention, such as selections of different transgenes and plasmids  
30 for the construction of the packaging cell lines and rAds, or selection or dosage of the viruses or immune modulators, are believed to be encompassed in the scope of the claims appended hereto.

## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

- (i) APPLICANT: Trustees of the University of Pennsylvania  
Wilson, James M.  
Fisher, Krishna J.  
Gao, Guang-Ping
- (ii) TITLE OF INVENTION: Recombinant Adenovirus and Adeno-  
Associated Virus, Cell Lines,  
and Methods of Production  
and Use Thereof
- (iii) NUMBER OF SEQUENCES: 5
- (iv) CORRESPONDENCE ADDRESS:
  - (A) ADDRESSEE: Howson and Howson
  - (B) STREET: Box 457, 321 Norristown Road
  - (C) CITY: Spring House
  - (D) STATE: PA
  - (E) COUNTRY: USA
  - (F) ZIP: 19477
- (v) COMPUTER READABLE FORM:
  - (A) MEDIUM TYPE: Floppy disk
  - (B) COMPUTER: IBM PC compatible
  - (C) OPERATING SYSTEM: PC-DOS/MS-DOS
  - (D) SOFTWARE: PatentIn Release 1.0 Version 1.30
- (vi) CURRENT APPLICATION DATA:
  - (A) APPLICATION NUMBER: WO
  - (B) FILING DATE:
  - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: US 08/462,014
  - (B) FILING DATE: 05-JUN-1995
- (vii) PRIOR APPLICATION DATA:
  - (A) APPLICATION NUMBER: US 08/549,489
  - (B) FILING DATE: 27-OCT-1995
- (viii) ATTORNEY/AGENT INFORMATION:
  - (A) NAME: Bak, Mary E.
  - (B) REGISTRATION NUMBER: 31,215
  - (C) REFERENCE/DOCKET NUMBER: GNVPN012CIPPCT
- (ix) TELECOMMUNICATION INFORMATION:
  - (A) TELEPHONE: (215) 540-9206
  - (B) TELEFAX: (215) 540-5818

## (2) INFORMATION FOR SEQ ID NO:1:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 3653 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: not relevant

## (ii) MOLECULE TYPE: cDNA

## (ix) FEATURE:

- (A) NAME/KEY: CDS
- (B) LOCATION: 1521..2405

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

CTGCATGTGT CAGAGGTTTT CACCGTCATC ACCGAAACGC GCGAGGCAGC	50
AAGCTTGGCA GAAATGGTTG AACTCCCGAG AGTGTCTTAC ACCTAGGGGA	100
GAAGCAGCCA AGGGGTTGTT TCCCACCAAG GACGACCCGT CTGCGCACAA	150
ACGGATGAGC CCATCAGACA AAGACATATT CATTCTCTGC TGCAAACCTG	200
GCATAGCTCT GCTTTGCCTG GGGCTATTGG GGAAGTTGC GGTTCGTGCT	250
CGCAGGGCTC TCACCCTTGA CTCTTTCAAT AATAACTCTT CTGTGCAAGA	300
TTACAATCTA AACAAATTCGG AGAACTCGAC CTTCTCCTG AGGCAAGGAC	350
CACAGCCAAC TTCCTCTTAC AAGCCGCATC GATTTTGTCC TTCAGAAATA	400
GAAATAAGAA TGCTTGCTAA AAATTATATT TTTACCAATA AGACCAATCC	450
AATAGGTAGA TTATTAGTTA CTATGTTAAG AAATGAATCA TTATCTTTTA	500
GTACTATTTT TACTCAAATT CAGAAGTTAG AAATGGGAAT AGAAAATAGA	550
AAGAGACGCT CAACCTCAAT TGAAGAACAG GTGCAAGGAC TATTGACCAC	600
AGGCCTAGAA GTAAAAAAGG GAAAAAAGAG TGTTTTTGTC AAAATAGGAG	650
ACAGGTGGTG GCAACCAGGG ACTTATAGGG GACCTTACAT CTACAGACCA	700
ACAGATGCCC CCTTACCATA TACAGGAAGA TATGACTTAA ATTGGGATAG	750
GTGGGTACCA GTCAATGGCT ATAAAGTGTT ATATAGATCC CTCCCCTTTC	800
GTGAAAGACT CGCCAGAGCT AGACCTCCTT GGTGTATGTT GTCTCAAGAA	850
AAGAAAGACG ACATGAAACA ACAGGTACAT GATTATATTT ATCTAGGAAC	900



AGGAATGCAC	TTTTGGGGAA	AGATTTTCCA	TACCAAGGAG	GGGACAGTGG	950
CTGGACTAAT	AGAACATTAT	TCTGCAAAAA	CTTATGGCAT	GAGTTATTAT	1000
GATTAGCCTT	GATTTGCCCA	ACCTTGCGGT	TCCCAAGGCT	TAAGTAAGTT	1050
TTTGGTTACA	AACTGTTCTT	AAAACAAGGA	TGTGAGACAA	GTGGTTTCCT	1100
GACTTGTTTT	GGTATCAAAG	GTTCTGATCT	GAGCTCTGAG	TGTTCTATTT	1150
TCCTATGTTT	TTTTGGAATT	TATCCAAATC	TTATGTAAAT	GCTTATGTAA	1200
ACCAAGATAT	AAAAGAGTGC	TGATTTTTTTG	AGTAAACTTG	CAACAGTCCT	1250
AACATTCACC	TCTTGTGTGT	TTGTGTCTGT	TCGCCATCCC	GTCTCCGCTC	1300
GTCACTTATC	CTTCACTTTC	CAGAGGGTCC	CCCCGCAGAC	CCCGGCGACC	1350
CTCAGGTCGG	CCGACTGCGG	CAGCTGGCGC	CCGAACAGGG	ACCCTCGGAT	1400
AAGTGACCCT	TGTCTTTATT	TCTACTATTT	TGTGTTTCGT	TTGTTTTGTC	1450
TCTATCTTGT	CTGGCTATCA	TCACAAGAGC	GGAACGGACT	CACCTCAGGG	1500
AACCAAGCTA	GCCCAATTCT	ATGACTACGT	CCGGCGTTCC	ATTTGGCATG	1550
AACTACGAC	CAACACGATC	TCGGTTGTCT	CGGCGCACTC	CGTACAGTAG	1600
GGATCGTCTA	CCTCCTTTTG	AGACAGAAAC	CCGCGCTACC	ATACTGGAGG	1650
ATCATCCGCT	GCTGCCCCGA	TGTAACACTT	TGACAATGCA	CAACGTGAGT	1700
TACGTGCGAG	GTCTTCCCTG	CAGTGTGGGA	TTTACGCTGA	TTCAGGAATG	1750
GGTTGTTCCC	TGGGATATGG	TTCTAACGCG	GGAGGAGCTT	GTAATCCTGA	1800
GGAAGTGTAT	GCACGTGTGC	CTGTGTTGTG	CCAACATTGA	TATCATGACG	1850
AGCATGATGA	TCCATGGTTA	CGAGTCCTGG	GCTCTCCACT	GTCATTGTTC	1900
CAGTCCCGGT	TCCCTGCAGT	GTATAGCCGG	CGGGCAGGTT	TTGGCCAGCT	1950
GGTTTAGGAT	GGTGGTGGAT	GGCGCCATGT	TTAATCAGAG	GTTTATATGG	2000
TACCGGGAGG	TGGTGAATTA	CAACATGCCA	AAAGAGGTAA	TGTTTATGTC	2050
CAGCGTGTTT	ATGAGGGGTC	GCCACTTAAT	CTACCTGCGC	TTGTGGTATG	2100
ATGGCCACGT	GGGTTCTGTG	GTCCCCGCCA	TGAGCTTTGG	ATACAGCGCC	2150
TTGCACTGTG	GGATTTTGAA	CAATATTGTG	GTGCTGTGCT	GCAGTTACTG	2200

TGCTGATTTA	AGTGAGATCA	GGGTGCGCTG	CTGTGCCCCG	AGGACAAGGC	2250
GCCTTATGCT	GCGGGCGGTG	CGAATCATCG	CTGAGGAGAC	CACTGCCATG	2300
TTGTATTTCCT	GCAGGACGGA	GCGGCGGCGG	CAGCAGTTTA	TTCGCGCGCT	2350
GCTGCAGCAC	CACCGCCCTA	TCCTGATGCA	CGATTATGAC	TCTACCCCCA	2400
TGTAGGGATC	CAAGCTTGCG	GGCGCATCGA	TGATATCAAG	CTTGCATGCC	2450
TGCAGGTCGA	CTCTAGAGGA	TCCCGGGTGG	NATCCCTGTG	ACCCCTCCCC	2500
AGTGCCTCTC	CTGGCCCTGG	AAGTTGGCAC	TCCAGTGCCC	ACCAGCCTTG	2550
TCCTAATAAA	ATTAAGTTGN	ATCATTTTGT	CTGACTAGGT	GTCCTTCTAT	2600
AATATTATGG	GGTGGAGGGG	GGTGGTATGG	AGCAANGGGN	AANTTGGNAA	2650
GACAANCTGT	AGGGCCTGCG	GGGTCTATTG	GGAACAAGCT	GGAGTGCAGT	2700
GGCACAATCT	TGGCTCACTG	CAATCTCCGC	CTCCTGGGTT	CAAGCGATTC	2750
TCCTGCCTCA	GACTCCCGAG	TTGTTGGGAT	TCCAGGCATG	CATGACCAGG	2800
CTCAGATAAT	TTTTGTTTTT	TTGGTAGAGA	CGGGGTTTCA	CCATATTGGN	2850
CAGGCTGGTC	TCCAACCTCCT	AATCTCAGGT	GATCTNCCCA	CCTTGGCCTC	2900
CCAAATTGCT	GGGATTACAG	GNGTGAACCA	CTGNTCCCTT	CCCTGTCCTT	2950
CTGATTTTAA	AATAACTATA	CCAGCAGGAG	GACGTCCAGA	CACAGCATAG	3000
GCTACCTGGC	CATGCCCAAC	CGGTGGGACA	TTTGAGTTGC	TTGCTTGGCA	3050
CTGTCCTCTC	ATGCGTTGGG	TCCACTCAGT	AGATGCCTGT	TGAATTGGGT	3100
ACGCGGCCAG	CTTGGCTGTG	GAATGTGTGT	CAGTTAGGGT	GTGGAAAGTC	3150
CCCAGGCTCC	CCAGCAGGCA	GAAGTATGCA	AAGCATGCAT	CTCAATTAGT	3200
CAGCAACCAG	GTGTGGAAAG	TCCCCAGGCT	CCCCAGCAGG	CAGAAGTATG	3250
CAAAGCATGC	ATCTCAATTA	GTCAGNAACC	ATAGNCCCGC	CCCTAACTCC	3300
GTCCATCCCG	GCCCTAACTC	NGGCCAGTTC	CGACCNNTNCT	CCGGCNNATG	3350
GNTGAGTAAT	TTGCNNGATT	TATGCAGNGG	GCGAGGNCGC	CTCGGGCTCT	3400
GAGNTNTTCC	AGAAGTAGTG	AGGAGGCTTT	NNTGGTGGAA	TTGATCAGCT	3450
TGGGATCTGA	TCAAGAGACA	GGATGAGGAT	CGNNNCGNAT	GATTGAACAA	3500

72

GATGGGTTGC ACGGAGGTTC TCCGNGCGCT TGGGTGGGGA GGNTATTCGG 3550  
 NTATTNTTGG TGNACAACAG NNAAACGGNT GTTCTGATGC CGCCGCGTTC 3600  
 NCGCTTTTCAG NGCAGGGGGG CCCCCCTTCT NTTGAGANNA GCNCCCCTTN 3650  
 TTG 3653

## (2) INFORMATION FOR SEQ ID NO:2:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 294 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS:
- (D) TOPOLOGY: not relevant

## (ii) MOLECULE TYPE: protein

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

Met	Thr	Thr	Ser	Gly	Val	Pro	Phe	Gly	Met	Thr	Leu	Arg	Pro	Thr	Arg	1	5	10	15
Ser	Arg	Leu	Ser	Arg	Arg	Thr	Pro	Tyr	Ser	Arg	Asp	Arg	Leu	Pro	Pro	20	25	30	
Phe	Glu	Thr	Glu	Thr	Arg	Ala	Thr	Ile	Leu	Glu	Asp	His	Pro	Leu	Leu	35	40	45	
Pro	Glu	Cys	Asn	Thr	Leu	Thr	Met	His	Asn	Val	Ser	Tyr	Val	Arg	Gly	50	55	60	
Leu	Pro	Cys	Ser	Val	Gly	Phe	Thr	Leu	Ile	Gln	Glu	Trp	Val	Val	Pro	65	70	75	80
Trp	Asp	Met	Val	Leu	Thr	Arg	Glu	Glu	Leu	Val	Ile	Leu	Arg	Lys	Cys	85	90	95	
Met	His	Val	Cys	Leu	Cys	Cys	Ala	Asn	Ile	Asp	Ile	Met	Thr	Ser	Met	100	105	110	
Met	Ile	Tyr	Gly	Tyr	Glu	Ser	Trp	Ala	Leu	His	Cys	His	Cys	Ser	Ser	115	120	125	
Pro	Gly	Ser	Leu	Gln	Cys	Ile	Ala	Gly	Gly	Gln	Val	Leu	Ala	Ser	Trp	130	135	140	
Phe	Arg	Met	Val	Val	Asp	Gly	Ala	Met	Phe	Asn	Gln	Arg	Phe	Ile	Trp	145	150	155	160

73

Tyr Arg Glu Val Val Asn Tyr Asn Met Pro Lys Glu Val Met Phe Met  
 165 170 175  
 Ser Ser Val Phe Met Arg Gly Arg His Leu Ile Tyr Leu Arg Tyr Trp  
 180 185 190  
 Tyr Asp Gly His Val Gly Ser Val Val Pro Ala Met Ser Phe Gly Tyr  
 195 200 205  
 Ser Ala Leu His Cys Gly Ile Leu Asn Asn Ile Val Val Leu Cys Cys  
 210 215 220  
 Ser Tyr Cys Ala Asp Leu Ser Glu Ile Arg Val Arg Cys Cys Ala Arg  
 225 230 235 240  
 Arg Thr Arg Arg Leu Met Leu Arg Ala Val Arg Ile Ile Ala Glu Glu  
 245 250 255  
 Thr Thr Ala Met Leu Tyr Ser Cys Arg Thr Glu Arg Arg Arg Gln Gln  
 260 265 270  
 Phe Ile Arg Ala Leu Leu Gln His His Arg Pro Ile Leu Met His Asp  
 275 280 285  
 Tyr Asp Ser Thr Pro Met  
 290

## (2) INFORMATION FOR SEQ ID NO:3:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 35408 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: not relevant
- (D) TOPOLOGY: not relevant

## (ii) MOLECULE TYPE: other nucleic acid

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

CATCATCAAT AATATACCTT ATTTTGGATT GAAGCCAATA TGATAATGAG 50  
 GGGGTGGAGT TTGTGACGTG GCGCGGGGCG TGGGAACGGG GCGGGTGACG 100  
 TAGTAGTGTG GCGGAAGTGT GATGTTGCAA GTGTGGCGGA ACACATGTAA 150  
 GCGACGGATG TGGCAAAAGT GACGTTTTTG GTGTGCGCCG GTGTACACAG 200  
 GAAGTGACAA TTTTCGCGCG GTTTTAGGCG GATGTTGTAG TAAATTTGGG 250  
 CGTAACCGAG TAAGATTTGG CCATTTTCGC GGGAAACTG AATAAGAGGA 300

AGTGAAATCT	GAATAATTTT	GTGTTACTCA	TAGCGCGTAA	TATTTGTCTA	350
GGGAGATCAG	CCTGCAGGTC	GTTACATAAC	TTACGGTAAA	TGGCCCGCCT	400
GGCTGACCGC	CCAACGACCC	CCGCCCATTG	ACGTCAATAA	TGACGTATGT	450
TCCCATAGTA	ACGCCAATAG	GGACTTTCCA	TTGACGTCAA	TGGGTGGAGT	500
ATTTACGGTA	AACTGCCCCAC	TTGGCAGTAC	ATCAAGTGTA	TCATATGCCA	550
AGTACGCCCC	CTATTGACGT	CAATGACGGT	AAATGGCCCCG	CCTGGCATT A	600
TGCCCAGTAC	ATGACCTTAT	GGGACTTTCC	TACTTGGCAG	TACATCTACT	650
CGAGGCCACG	TTCTGCTTCA	CTCTCCCCAT	CTCCCCCCCC	TCCCCACCCC	700
CAATTTTGTA	TTTATTTATT	TTTTAATTAT	TTTGTGCAGC	GATGGGGGCG	750
GGGGGGGGGG	GGGGGCGCGC	GCCAGGCGGG	GCGGGGCGGG	GCGAGGGGCG	800
GGGCGGGGCG	AGGCGGAGAG	GTGCGGCGGC	AGCCAATCAG	AGCGGCGCGC	850
TCCGAAAGTT	TCCTTTTATG	GCGAGGCGGC	GGCGGCGGCG	GCCCTATAAA	900
AAGCGAAGCG	CGCGGCGGGC	GGGAGCGGGA	TCAGCCACCG	CGGTGGCGGC	950
CGCAATTCCC	GGGGATCGAA	AGAGCCTGCT	AAAGCAAAAA	AGAAGTCACC	1000
ATGTCGTTTA	CTTTGACCAA	CAAGAACGTG	ATTTTCGTTG	CCGGTCTGGG	1050
AGGCATTGGT	CTGGACACCA	GCAAGGAGCT	GCTCAAGCGC	GATCCCGTCG	1100
TTTTACAACG	TCGTGACTGG	GAAAACCCTG	GCGTTACCCA	ACTTAATCGC	1150
CTTGCAGCAC	ATCCCCCTTT	CGCCAGCTGG	CGTAATAGCG	AAGAGGCCCC	1200
CACCGATCGC	CCTTCCCAAC	AGTTGCGCAG	CCTGAATGGC	GAATGGCGCT	1250
TTGCCTGGTT	TCCGGCACCA	GAAGCGGTGC	CGGAAAGCTG	GCTGGAGTGC	1300
GATCTTCCTG	AGGCCGATAC	TGTCGTCGTC	CCCTCAAAC T	GGCAGATGCA	1350
CGGTTACGAT	GCGCCCATCT	ACACCAACGT	AACCTATCCC	ATTACGGTCA	1400
ATCCGCCGTT	TGTTCCACAG	GAGAATCCGA	CGGGTTGTTA	CTCGCTCACA	1450
TTTAATGTTG	ATGAAAGCTG	GCTACAGGAA	GGCCAGACGC	GAATTATTTT	1500
TGATGGCGTT	AACTCGGCGT	TTCATCTGTG	GTGCAACGGG	CGCTGGGTCC	1550
GTTACGGCCA	GGACAGTCGT	TTGCCGTCTG	AATTTGACCT	GAGCGCATTT	1600

TTACGCGCCG	GAGAAAACCG	CCTCGCGGTG	ATGGTGCTGC	GTTGGAGTGA	1650
CGGCAGTTAT	CTGGAAGATC	AGGATATGTG	GCGGATGAGC	GGCATTTTCC	1700
GTGACGTCTC	GTTGCTGCAT	AAACCGACTA	CACAAATCAG	CGATTTCCAT	1750
GTTGCCACTC	GCTTTAATGA	TGATTTCAGC	CGCGCTGTAC	TGGAGGCTGA	1800
AGTTCAGATG	TGCGGCGAGT	TGCGTGACTA	CCTACGGGTA	ACAGTTTCTT	1850
TATGGCAGGG	TGAAACGCAG	GTCGCCAGCG	GCACCGCGCC	TTTCGGCGGT	1900
GAAATTATCG	ATGAGCGTGG	TGGTTATGCC	GATCGCGTCA	CACTACGTCT	1950
GAACGTCGAA	AACCCGAAAC	TGTGGAGCGC	CGAAATCCCG	AATCTCTATC	2000
GTGCGGTGGT	TGAACTGCAC	ACCGCCGACG	GCACGCTGAT	TGAAGCAGAA	2050
GCCTGCGATG	TCGGTTTCCG	CGAGGTGCGG	ATTGAAAATG	GTCTGCTGCT	2100
GCTGAACGGC	AAGCCGTTGC	TGATTCGAGG	CGTTAACCGT	CACGAGCATC	2150
ATCCTCTGCA	TGGTCAGGTC	ATGGATGAGC	AGACGATGGT	GCAGGATATC	2200
CTGCTGATGA	AGCAGAACAA	CTTTAACGCC	GTGCGCTGTT	CGCATTATCC	2250
GAACCATCCG	CTGTGGTACA	CGCTGTGCGA	CCGCTACGGC	CTGTATGTGG	2300
TGGATGAAGC	CAATATTGAA	ACCCACGGCA	TGGTGCCAAT	GAATCGTCTG	2350
ACCGATGATC	CGCGCTGGCT	ACCGGCGATG	AGCGAACGCG	TAACGCGAAT	2400
GGTGCAGCGC	GATCGTAATC	ACCCGAGTGT	GATCATCTGG	TCGCTGGGGA	2450
ATGAATCAGG	CCACGGCGCT	AATCACGACG	CGCTGTATCG	CTGGATCAAA	2500
TCTGTCGATC	CTTCCCGCCC	GGTGCAGTAT	GAAGGCGGCG	GAGCCGACAC	2550
CACGGCCACC	GATATTATTT	GCCCGATGTA	CGCGCGCGTG	GATGAAGACC	2600
AGCCCTTCCC	GGCTGTGCCG	AAATGGTCCA	TCAAAAAATG	GCTTTCGCTA	2650
CCTGGAGAGA	CGCGCCCGCT	GATCCTTTGC	GAATACGCCC	ACGCGATGGG	2700
TAACAGTCTT	GGCGGTTTCG	CTAAATACTG	GCAGGCGTTT	CGTCAGTATC	2750
CCCGTTTACA	GGGCGGCTTC	GTCTGGGACT	GGGTGGATCA	GTCGCTGATT	2800
AAATATGATG	AAAACGGCAA	CCCGTGGTCTG	GCTTACGGCG	GTGATTTTGG	2850
CGATACGCCG	AACGATCGCC	AGTTCTGTAT	GAACGGTCTG	GTCTTTGCCG	2900

ACCGCACGCC	GCATCCAGCG	CTGACGGAAG	CAAAACACCA	GCAGCAGTTT	2950
TTCCAGTTCC	GTTTATCCGG	GCAAACCATC	GAAGTGACCA	GCGAATACCT	3000
GTTCCGTCAT	AGCGATAACG	AGCTCCTGCA	CTGGATGGTG	GCGCTGGATG	3050
GTAAGCCGCT	GGCAAGCGGT	GAAGTGCCTC	TGGATGTCGC	TCCACAAGGT	3100
AAACAGTTGA	TTGAACTGCC	TGAACTACCG	CAGCCGGAGA	GCGCCGGGCA	3150
ACTCTGGCTC	ACAGTACGCG	TAGTGCAACC	GAACGCGACC	GCATGGTCAG	3200
AAGCCGGGCA	CATCAGCGCC	TGGCAGCAGT	GGCGTCTGGC	GGAAAACCTC	3250
AGTGTGACGC	TCCCCGCCGC	GTCCCACGCC	ATCCCCGCATC	TGACCACCAG	3300
CGAAATGGAT	TTTTGTCATCG	AGCTGGGTAA	TAAGCGTTGG	CAATTTAACC	3350
GCCAGTCAGG	CTTTCTTTCA	CAGATGTGGA	TTGGCGATAA	AAAACAAC TG	3400
CTGACGCCGC	TGCGCGATCA	GTTCACCCGT	GCACCGCTGG	ATAACGACAT	3450
TGGCGTAAAGT	GAAGCGACCC	GCATTGACCC	TAACGCCTGG	GTCGAACGCT	3500
GGAAGGCGGC	GGGCCATTAC	CAGGCCGAAG	CAGCGTTGTT	GCAGTGCACG	3550
GCAGATACAC	TTGCTGATGC	GGTGCTGATT	ACGACCGCTC	ACGCGTGGCA	3600
GCATCAGGGG	AAAACCTTAT	TTATCAGCCG	GAAAACCTAC	CGGATTGATG	3650
GTAGTGGTCA	AATGGCGATT	ACCGTTGATG	TTGAAGTGGC	GAGCGATACA	3700
CCGCATCCGG	CGCGGATTGG	CCTGAACTGC	CAGCTGGCGC	AGGTAGCAGA	3750
GCGGGTAAAC	TGGCTCGGAT	TAGGGCCGCA	AGAAAACCTAT	CCCGACCGCC	3800
TTACTGCCGC	CTGTTTTGAC	CGCTGGGATC	TGCCATTGTC	AGACATGTAT	3850
ACCCCGTACG	TCTTCCCGAG	CGAAAACGGT	CTGCGCTGCG	GGACGCGCGA	3900
ATTGAATTAT	GGCCACACAC	AGTGGCGCGG	CGACTTCCAG	TTCAACATCA	3950
GCCGCTACAG	TCAACAGCAA	CTGATGGAAA	CCAGCCATCG	CCATCTGCTG	4000
CACGCGGAAG	AAGGCACATG	GCTGAATATC	GACGGTTTCC	ATATGGGGAT	4050
TGGTGGCGAC	GACTCCTGGA	GCCCGTCAGT	ATCGGCGGAA	TTACAGCTGA	4100
GCGCCGGTCG	CTACCATTAC	CAGTTGGTCT	GGTGTCAAAA	ATAATAATAA	4150
CCGGGCAGGC	CATGTCTGCC	CGTATTTTCGC	GTAAGGAAAT	CCATTATGTA	4200

CTATTTAAAA	AACACAAACT	TTTGGATGTT	CGGTTTATTC	TTTTTCTTTT	4250
ACTTTTTTAT	CATGGGAGCC	TACTTCCCGT	TTTTCCCGAT	TTGGCTACAT	4300
GACATCAACC	ATATCAGCAA	AAGTGATACG	GGTATTATTT	TTGCCGCTAT	4350
TTCTCTGTTC	TCGCTATTAT	TCCAACCGCT	GTTTGGTCTG	CTTCTGACA	4400
AACTCGGCCT	CGACTCTAGG	CGGCCGCGGG	GATCCAGACA	TGATAAGATA	4450
CATTGATGAG	TTTGGACAAA	CCACAAC TAG	AATGCAGTGA	AAAAAATGCT	4500
TTATTTGTGA	AATTTGTGAT	GCTATTGCTT	TATTTGTAAC	CATTATAAGC	4550
TGCAATAAAC	AAGTTAACAA	CAACAATTGC	ATTCATTTTA	TGTTTCAGGT	4600
TCAGGGGGAG	GTGTGGGAGG	TTTTTTCGGA	TCCTCTAGAG	TCGACCTGCA	4650
GGCTGATCAG	TGGAAGGTGC	TGAGGTACGA	TGAGACCCGC	ACCAGGTGCA	4700
GACCTGCGA	GTGTGGCGGT	AAACATATTA	GGAACCAGCC	TGTGATGCTG	4750
GATGTGACCG	AGGAGCTGAG	GCCCGATCAC	TTGGTGCTGG	CCTGCACCCG	4800
CGCTGAGTTT	GGCTCTAGCG	ATGAAGATAC	AGATTGAGGT	ACTGAAATGT	4850
GTGGGCCGTG	CTTAAGGGTG	GGAAAGAATA	TATAAGGTGG	GGGTCTTATG	4900
TAGTTTTGTA	TCTGTTTTGC	AGCAGCCGCC	GCCGCCATGA	GCACCAACTC	4950
GTTTGATGGA	AGCATTGTGA	GCTCATATTT	GACAACGCGC	ATGCCCCCAT	5000
GGGCCGGGGT	GCGTCAGAAT	GTGATGGGCT	CCAGCATTGA	TGGTCGCCCC	5050
GTCCTGCCCC	CAAAC TCTAC	TACCTTGACC	TACGAGACCG	TGTCTGGAAC	5100
GCCGTTGGAG	ACTGCAGCCT	CCGCCGCCGC	TTCAGCCGCT	GCAGCCACCG	5150
CCCGCGGGAT	TGTGACTGAC	TTTGCTTTCC	TGAGCCCGCT	TGCAAGCAGT	5200
GCAGCTTCCC	GTTCATCCGC	CCGCGATGAC	AAGTTGACGG	CTCTTTTGGC	5250
ACAATTGGAT	TCTTTGACCC	GGGAAC TTAA	TGTCGTTTCT	CAGCAGCTGT	5300
TGGATCTGCG	CCAGCAGGTT	TCTGCCCTGA	AGGCTTCCTC	CCCTCCCAAT	5350
GCGGTTTAAA	ACATAAATAA	AAAACCAGAC	TCTGTTTGGA	TTTGGATCAA	5400
GCAAGTGTCT	TGCTGTCTTT	ATTTAGGGGT	TTTGCGCGCG	CGGTAGGCCC	5450
GGGACCAGCG	GTCTCGGTCG	TTGAGGGTCC	TGTGTATTTT	TTCCAGGACG	5500



TGGTAAAGGT	GACTCTGGAT	G TTCAGATAC	ATGGGCATAA	GCCCGTCTCT	5550
GGGGTGGAGG	TAGCACC ACT	GCAGAGCTTC	ATGCTGCGGG	GTGGTGTTGT	5600
AGATGATCCA	GTCGTAGCAG	GAGCGCTGGG	CGTGGTGCCT	AAAAATGTCT	5650
TTCAGTAGCA	AGCTGATTGC	CAGGGGCAGG	CCCTTGGTGT	AAGTGTTTAC	5700
AAAGCGGTTA	AGCTGGGATG	GGTG CATACG	TGGGGATATG	AGATGCATCT	5750
TGGACTGTAT	TTTTAGGTTG	GCTATGTTCC	CAGCCATATC	CCTCCGGGGA	5800
TTCATGTTGT	GCAGAACCAC	CAGCACAGTG	TATCCGGTGC	ACTTGGGAAA	5850
TTTGTCATGT	AGCTTAGAAG	GAAATGCGTG	GAAGA ACTTG	GAGACGCCCT	5900
TGTGACCTCC	AAGATTTTCC	ATGCATT CGT	CCATAATGAT	GGCAATGGGC	5950
CCACGGGCGG	CGGCCTGGGC	GAAGATATTT	CTGGGATCAC	TAACGTCATA	6000
GTTGTGTTCC	AGGATGAGAT	CGTCATAGGC	CATTTTTTACA	AAGCGCGGGC	6050
GGAGGGTGCC	AGACTGCGGT	ATAATGGTTC	CATCCGGCCC	AGGGGCGTAG	6100
TTACCCTCAC	AGATTTGCAT	TTCCCACGCT	TTGAGTTCAG	ATGGGGGGAT	6150
CATGTCTACC	TGCGGGGCGA	TGAAGAAAAC	GGTTTCCGGG	G TAGGGGAGA	6200
TCAGCTGGGA	AGAAAGCAGG	TTCTTGAGCA	GCTGCGACTT	ACCGCAGCCG	6250
GTGGGCCCCG	AAATCACACC	TATTACCGGG	TGCAACTGGT	AGTTAAGAGA	6300
GCTGCAGCTG	CCGTCATCCC	TGAGCAGGGG	GGCCACTTCG	TTAAGCATGT	6350
CCCTGACTCG	CATGTTTTCC	CTGACCAAAT	CCGCCAGAAG	GCGCTCGCCG	6400
CCCAGCGATA	GCAGTTCCTG	CAAGGAAGCA	AAGTTTTTCA	ACGGTTTGAG	6450
ACCGTCCGCC	G TAGGCATGC	TTTTGAGCGT	TTGACCAAGC	AGTTCCAGGC	6500
GGTCCCACAG	CTCGGTCACC	TGCTCTACGG	CATCTCGATC	CAGCATATCT	6550
CCTCGTTTCG	CGGGTTGGGG	CGGCTTTCGC	TGTACGGCAG	TAGTCGGTGC	6600
TCGTCCAGAC	GGGCCAGGGT	CATGTCTTTC	CACGGGCGCA	GGGTCCCTCGT	6650
CAGCGTAGTC	TGGGTCACGG	TGAAGGGGTG	CGCTCCGGGC	TGCGCGCTGG	6700
CCAGGGTGCG	CTTGAGGCTG	GTCCTGCTGG	TGCTGAAGCG	CTGCCGGTCT	6750
TCGCCCTGCG	CGTCGGCCAG	G TAGCATTTG	ACCATGGTGT	CATAGTCCAG	6800

CCCCTCCGCG	GCGTGGCCCT	TGGCGCGCAG	CTTGCCCTTG	GAGGAGGCGC	6850
CGCACGAGGG	GCAGTGCAGA	CTTTTGAGGG	CGTAGAGCTT	GGGCGCGAGA	6900
AATACCGATT	CCGGGGAGTA	GGCATCCGCG	CCGCAGGCCC	CGCAGACGGT	6950
CTCGCATTCC	ACGAGCCAGG	TGAGCTCTGG	CCGTTCGGGG	TCAAAAACCA	7000
GGTTTCCCCC	ATGCTTTTTG	ATGCGTTTCT	TACCTCTGGT	TTCCATGAGC	7050
CGGTGTCCAC	GCTCGGTGAC	GAAAAGGCTG	TCCGTGTCCC	CGTATACAGA	7100
CTTGAGAGGC	CTGTCCTCGA	GCGGTGTTCC	GCGGTCCTCC	TCGTATAGAA	7150
ACTCGGACCA	CTCTGAGACA	AAGGCTCGCG	TCCAGGCCAG	CACGAAGGAG	7200
GCTAAGTGGG	AGGGGTAGCG	GTCGTTGTCC	ACTAGGGGGT	CCACTCGCTC	7250
CAGGGTGTGA	AGACACATGT	CGCCCTCTTC	GGCATCAAGG	AAGGTGATTG	7300
GTTTGTAGGT	GTAGGCCACG	TGACCGGGTG	TTCCTGAAGG	GGGGCTATAA	7350
AAGGGGGTGG	GGGCGCGTTC	GTCCTCACTC	TCTTCCGCAT	CGCTGTCTGC	7400
GAGGGCCAGC	TGTTGGGGTG	AGTACTCCCT	CTGAAAAGCG	GGCATGACTT	7450
CTGCGCTAAG	ATTGTCAGTT	TCCAAAAACG	AGGAGGATTT	GATATTCACC	7500
TGGCCCGCGG	TGATGCCTTT	GAGGGTGGCC	GCATCCATCT	GGTCAGAAAA	7550
GACAATCTTT	TTGTTGTCAA	GCTTGGTGGC	AAACGACCCG	TAGAGGGCGT	7600
TGGACAGCAA	CTTGGCGATG	GAGCGCAGGG	TTTGGTTTTT	GTCGCGATCG	7650
GCGCGCTCCT	TGGCCGCGAT	GTTTAGCTGC	ACGTATTTCG	GCGCAACGCA	7700
CCGCCATTCT	GGAAAGACGG	TGGTGCGCTC	GTCGGGCACC	AGGTGCACGC	7750
GCCAACCGCG	GTTGTGCAGG	GTGACAAGGT	CAACGCTGGT	GGCTACCTCT	7800
CCGCGTAGGC	GCTCGTTGGT	CCAGCAGAGG	CGGCCGCCCT	TGCGCGAGCA	7850
GAATGGCGGT	AGGGGGTCTA	GCTGCGTCTC	GTCCGGGGGG	TCTGCGTCCA	7900
CGGTAAAGAC	CCCGGGCAGC	AGGCGCGCGT	CGAAGTAGTC	TATCTTGCAT	7950
CCTTGCAAGT	CTAGCGCCTG	CTGCCATGCG	CGGGCGGCAA	GCGCGCGCTC	8000
GTATGGGTTG	AGTGGGGGAC	CCCATGGCAT	GGGGTGGGTG	AGCGCGGAGG	8050
CGTACATGCC	GCAAATGTCT	TAAACGTAGA	GGGGCTCTCT	GAGTATTCCA	8100

80

AGATATGTAG	GGTAGCATCT	TCCACCGCGG	ATGCTGGCGC	GCACGTAATC	8150
GTATAGTTTCG	TGCGAGGGAG	CGAGGAGGTC	GGGACCGAGG	TTGCTACGGG	8200
CGGGCTGCTC	TGCTCGGAAG	ACTATCTGCC	TGAAGATGGC	ATGTGAGTTG	8250
GATGATATGG	TTGGACGCTG	GAAGACGTTG	AAGCTGGCGT	CTGTGAGACC	8300
TACCGCGTCA	CGCACGAAGG	AGGCGTAGGA	GTCGCGCAGC	TTGTTGACCA	8350
GCTCGGCGGT	GACCTGCACG	TCTAGGGCGC	AGTAGTCCAG	GGTTTCCTTG	8400
ATGATGTCAT	ACTTATCCTG	TCCCTTTTTT	TTCCACAGCT	CGCGGTTGAG	8450
GACAAACTCT	TCGCGGTCTT	TCCAGTACTC	TTGGATCGGA	AACCCGTCGG	8500
CCTCCGAACG	GTAAGAGCCT	AGCATGTAGA	ACTGGTTGAC	GGCCTGGTAG	8550
GCGCAGCATC	CCTTTTCTAC	GGGTAGCGCG	TATGCCTGCG	CGGCCTTCCG	8600
GAGCGAGGTG	TGGGTGAGCG	CAAAGGTGTC	CCTGACCATG	ACTTTGAGGT	8650
ACTGGTATTT	GAAGTCAGTG	TCGTCGCATC	CGCCCTGCTC	CCAGAGCAAA	8700
AAGTCCGTGC	GCTTTTTTGA	ACGCGGATTT	GGCAGGGCGA	AGGTGACATC	8750
GTTGAAGAGT	ATCTTTCCCG	CGCGAGGCAT	AAAGTTGCGT	GTGATGCGGA	8800
AGGGTCCCGG	CACCTCGGAA	CGGTTGTTAA	TTACCTGGGC	GGCGAGCACG	8850
ATCTCGTCAA	AGCCGTTGAT	GTTGTGGCCC	ACAATGTAAA	GTTCCAAGAA	8900
GCGCGGGATG	CCCTTGATGG	AAGGCAATTT	TTTAAGTTCC	TCGTAGGTGA	8950
GCTCTTCAGG	GGAGCTGAGC	CCGTGCTCTG	AAAGGGCCCA	GTCTGCAAGA	9000
TGAGGGTTGG	AAGCGACGAA	TGAGCTCCAC	AGGTCACGGG	CCATTAGCAT	9050
TTGCAGGTGG	TCGCGAAAGG	TCCTAAACTG	GCGACCTATG	GCCATTTTTT	9100
CTGGGGTGAT	GCAGTAGAAG	GTAAGCGGGT	CTTGTTCCCA	GCGGTCCCAT	9150
CCAAGGTTCG	CGGCTAGGTC	TCGCGCGGCA	GTCAC TAGAG	GCTCATCTCC	9200
GCCGAAC TTC	ATGACCAGCA	TGAAGGGCAC	GAGCTGCTTC	CCAAAGGCCC	9250
CCATCCAAGT	ATAGGTCTCT	ACATCGTAGG	TGACAAAGAG	ACGCTCGGTG	9300
CGAGGATGCG	AGCCGATCGG	GAAGAACTGG	ATCTCCCGCC	ACCAATTGGA	9350
GGAGTGGCTA	TTGATGTGGT	GAAAGTAGAA	GTCCCTGCGA	CGGGCCGAAC	9400

ACTCGTGCTG	GCTTTTGTA	AAACGTGCGC	AGTACTGGCA	GCGGTGCACG	9450
GGCTGTACAT	CCTGCACGAG	GTTGACCTGA	CGACCGCGCA	CAAGGAAGCA	9500
GAGTGGGAAT	TTGAGCCCCT	CGCCTGGCGG	GTTTGGCTGG	TGGTCTTCTA	9550
CTTCGGCTGC	TTGTCCTTGA	CCGTCTGGCT	GCTCGAGGGG	AGTTACGGTG	9600
GATCGGACCA	CCACGCCGCG	CGAGCCCAAA	GTCCAGATGT	CCGCGCGCGG	9650
CGGTGCGGAGC	TTGATGACAA	CATCGCGCAG	ATGGGAGCTG	TCCATGGTCT	9700
GGAGCTCCCG	CGGCGTCAGG	TCAGGCGGGA	GCTCCTGCAG	GTTTACCTCG	9750
CATAGACGGG	TCAGGGCGCG	GGCTAGATCC	AGGTGATACC	TAATTTCCAG	9800
GGGCTGGTTG	GTGGCGGCGT	CGATGGCTTG	CAAGAGGCCG	CATCCCCGCG	9850
GCGCGACTAC	GGTACCGCGC	GGCGGGCGGT	GGGCCGCGGG	GGTGTCTTTG	9900
GATGATGCAT	CTAAAAGCGG	TGACGCGGGC	GAGCCCCCGG	AGGTAGGGGG	9950
GGCTCCGGAC	CCGCCGGGAG	AGGGGGCAGG	GGCACGTCGG	CGCCGCGCGC	10000
GGGCAGGAGC	TGGTGCTGCG	CGCGTAGGTT	GCTGGCGAAC	GCGACGACGC	10050
GGCGGTTGAT	CTCCTGAATC	TGGCGCCTCT	GCGTGAAGAC	GACGGGCCCCG	10100
GTGAGCTTGA	GCCTGAAAGA	GAGTTCGACA	GAATCAATTT	CGGTGTCGTT	10150
GACGGCGGCC	TGGCGCAAAA	TCTCCTGCAC	GTCTCCTGAG	TTGTCTTGAT	10200
AGGCGATCTC	GGCCATGAAC	TGCTCGATCT	CTTCCTCCTG	GAGATCTCCG	10250
CGTCCGGCTC	GCTCCACGGT	GGCGGCGAGG	TCGTTGGAAA	TGCGGGCCAT	10300
GAGCTGCGAG	AAGGCGTTGA	GGCCTCCCTC	GTTCCAGACG	CGGCTGTAGA	10350
CCACGCCCCC	TTCGGCATCG	CGGGCGCGCA	TGACCACCTG	CGCGAGATTG	10400
AGCTCCACGT	GCCGGGCGAA	GACGGCGTAG	TTTCGCAGGC	GCTGAAAGAG	10450
GTAGTTGAGG	GTGGTGGCGG	TGTGTTCTGC	CACGAAGAAG	TACATAACCC	10500
AGCGTCGCAA	CGTGGATTCTG	TTGATATCCC	CCAAGGCCTC	AAGGCGCTCC	10550
ATGGCCTCGT	AGAAGTCCAC	GGCGAAGTTG	AAAACTGGG	AGTTGCGCGC	10600
CGACACGGTT	AACTCCTCCT	CCAGAAGACG	GATGAGCTCG	GCGACAGTGT	10650
CGCGCACCTC	GCGCTCAAAG	GCTACAGGGG	CCTCTTCTTC	TTCTTCAATC	10700

TCCTCTTCCA	TAAGGGCCTC	CCCTTCTTCT	TCTTCTGGCG	GCGGTGGGGG	10750
AGGGGGGACA	CGGCGGCGAC	GACGGCGCAC	CGGGAGGCGG	TCGACAAAGC	10800
GCTCGATCAT	CTCCCCGCGG	CGACGGCGCA	TGGTCTCGGT	GACGGCGCGG	10850
CCGTTCTCGC	GGGGGCGCAG	TTGGAAGACG	CCGCCCCTCA	TGTCCCGGTT	10900
ATGGGTGTC	GGGGGGCTGC	CATGCGGCAG	GGATACGGCG	CTAACGATGC	10950
ATCTCAACAA	TTGTTGTGTA	GGTACTCCGC	CGCCGAGGGA	CCTGAGCGAG	11000
TCCGCATCGA	CCGGATCGGA	AAACCTCTCG	AGAAAGGCGT	CTAACCAGTC	11050
ACAGTCGCAA	GGTAGGCTGA	GCACCGTGGC	GGGCGGCAGC	GGGCGGCGGT	11100
CGGGGTTGTT	TCTGGCGGAG	GTGCTGCTGA	TGATGTAATT	AAAGTAGGCG	11150
GTCTTGAGAC	GGCGGATGGT	CGACAGAAGC	ACCATGTCCT	TGGGTCCGGC	11200
CTGCTGAATG	CGCAGGCGGT	CGGCCATGCC	CCAGGCTTCG	TTTGTACATC	11250
GGCGCAGGTC	TTTGTAGTAG	TCTTGCAATG	GCCTTTCTAC	CGGCACCTCT	11300
TCTTCTCCTT	CCTCTTGTCC	TGCATCTCTT	GCATCTATCG	CTGCGGCGGC	11350
GGCGGAGTTT	GGCCGTAGGT	GGCGCCCTCT	TCCTCCCATG	CGTGTGACCC	11400
CGAAGCCCCT	CATCGGCTGA	AGCAGGGCTA	GGTCGGCGAC	AACGCGCTCG	11450
GCTAATATGG	CCTGCTGCAC	CTGCGTGAGG	GTAGACTGGA	AGTCATCCAT	11500
GTCCACAAAG	CGGTGGTATG	CGCCCGTGTT	GATGGTGTA	GTGCAGTTGG	11550
CCATAACGGA	CCAGTTAACG	GTCTGGTGAC	CCGGCTGCGA	GAGCTCGGTG	11600
TACCTGAGAC	GCGAGTAAGC	CCTCGAGTCA	AATACGTAGT	CGTTGCAAGT	11650
CCGCACCAGG	TACTGGTATC	CCACCAAAAA	GTGCGGCGGC	GGCTGGCGGT	11700
AGAGGGGCCA	GCGTAGGGTG	GCCGGGGCTC	CGGGGGCGAG	ATCTTCCAAC	11750
ATAAGGCGAT	GATATCCGTA	GATGTACCTG	GACATCCAGG	TGATGCCGGC	11800
GGCGGTGGTG	GAGGCGCGCG	GAAAGTCGCG	GACGCGGTTC	CAGATGTTGC	11850
GCAGCGGCAA	AAAGTGCTCC	ATGGTCGGGA	CGCTCTGGCC	GGTCAGGCGC	11900
GCGCAATCGT	TGACGCTCTA	GACCGTGCAA	AAGGAGAGCC	TGTAAGCGGG	11950
CACTCTTCCG	TGGTCTGGTG	GATAAATTCG	CAAGGGTATC	ATGGCGGACG	12000

ACCGGGGTTT	GAGCCCCGTA	TCCGGCCGTC	CGCCGTGATC	CATGCGGTTA	12050
CCGCCCCGCT	GTCGAACCCA	GGTGTGCGAC	GTCAGACAAC	GGGGGAGTGC	12100
TCCTTTTGGC	TTCCTTCCAG	GCGCGGCGGC	TGCTGCGCTA	GCTTTTTTGG	12150
CCACTGGCCG	CGCGCAGCGT	AAGCGGTTAG	GCTGGAAAGC	GAAAGCATTA	12200
AGTGGCTCGC	TCCCTGTAGC	CGGAGGGTTA	TTTTCCAAGG	GTTGAGTCGC	12250
GGGACCCCCG	GTTTCGAGTCT	CGGACCGGCC	GGACTGCGGC	GAACGGGGGT	12300
TTGCCTCCCC	GTCATGCAAG	ACCCCGCTTG	CAAATTCCTC	CGGAAACAGG	12350
GACGAGCCCC	TTTTTTGCTT	TTCCCAGATG	CATCCGGTGC	TGCGGCAGAT	12400
GCGCCCCCCT	CCTCAGCAGC	GGCAAGAGCA	AGAGCAGCGG	CAGACATGCA	12450
GGGCACCCTC	CCCTCCTCCT	ACCGCGTCAG	GAGGGGCGAC	ATCCGCGGTT	12500
GACGCGGCAG	CAGATGGTGA	TTACGAACCC	CCGCGGCGCC	GGGCCC GGCA	12550
CTACCTGGAC	TTGGAGGAGG	GCGAGGGCCT	GGCGCGGCTA	GGAGCGCCCT	12600
CTCCTGAGCG	GTACCCAAGG	GTGCAGCTGA	AGCGTGATAC	GCGTGAGGCG	12650
TACGTGCCGC	GGCAGAACCT	GTTTCGCGAC	CGCGAGGGAG	AGGAGCCCGA	12700
GGAGATGCGG	GATCGAAAGT	TCCACGCAGG	GCGCGAGCTG	CGGCATGGCC	12750
TGAATCGCGA	GCGGTTGCTG	CGCGAGGAGG	ACTTTGAGCC	CGACGCGCGA	12800
ACCGGGATTA	GTCCCGCGCG	CGCACACGTG	GCGGCCGCCG	ACCTGGTAAC	12850
CGCATACGAG	CAGACGGTGA	ACCAGGAGAT	TAACTTTCAA	AAAAGCTTTA	12900
ACAACCACGT	GCGTACGCTT	GTGGCGCGCG	AGGAGGTGGC	TATAGGACTG	12950
ATGCATCTGT	GGGACTTTGT	AAGCGCGCTG	GAGCAAAACC	CAAATAGCAA	13000
GCCGCTCATG	GCGCAGCTGT	TCCTTATAGT	GCAGCACAGC	AGGGACAACG	13050
AGGCATTTCAG	GGATGCGCTG	CTAAACATAG	TAGAGCCCGA	GGGCCGCTGG	13100
CTGCTCGATT	TGATAAACAT	CCTGCAGAGC	ATAGTGGTGC	AGGAGCGCAG	13150
CTTGAGCCTG	GCTGACAAGG	TGGCCGCCAT	CAACTATTCC	ATGCTTAGCC	13200
TGGGCAAGTT	TTACGCCCCG	AAGATATACC	ATACCCCTTA	CGTTCCCATATA	13250
GACAAGGAGG	TAAAGATCGA	GGGGTTCTAC	ATGCGCATGG	CGCTGAAGGT	13300

GCTTACCTTG	AGCGACGACC	TGGGCGTTTA	TCGCAACGAG	CGCATCCACA	13350
AGGCCGTGAG	CGTGAGCCGG	CGGCGCGAGC	TCAGCGACCG	CGAGCTGATG	13400
CACAGCCTGC	AAAGGGCCCT	GGCTGGCACG	GGCAGCGGCG	ATAGAGAGGC	13450
CGAGTCCTAC	TTTGACGCGG	GCGCTGACCT	GCGCTGGGCC	CCAAGCCGAC	13500
GCGCCCTGGA	GGCAGCTGGG	GCCGGACCTG	GGCTGGCGGT	GGCACCCGCG	13550
CGCGCTGGCA	ACGTCGGCGG	CGTGGAGGAA	TATGACGAGG	ACGATGAGTA	13600
CGAGCCAGAG	GACGGCGAGT	ACTAAGCGGT	GATGTTTCTG	ATCAGATGAT	13650
GCAAGACGCA	ACGGACCCGG	CGGTGCGGGC	GGCGCTGCAG	AGCCAGCCGT	13700
CCGGCCTTAA	CTCCACGGAC	GACTGGCGCC	AGGTCATGGA	CCGCATCATG	13750
TCGCTGACTG	CGCGCAATCC	TGACGCGTTC	CGGCAGCAGC	CGCAGGCCAA	13800
CCGGCTCTCC	GCAATTCTGG	AAGCGGTGGT	CCCGGCGCGC	GCAAACCCCA	13850
CGCACGAGAA	GGTGCTGGCG	ATCGTAAACG	CGCTGGCCGA	AAACAGGGCC	13900
ATCCGGCCCG	ACGAGGCCGG	CCTGGTCTAC	GACGCGCTGC	TTCAGCGCGT	13950
GGCTCGTTAC	AACAGCGGCA	ACGTGCAGAC	CAACCTGGAC	CGGCTGGTGG	14000
GGGATGTGCG	CGAGGCCGTG	GCGCAGCGTG	AGCGCGCGCA	GCAGCAGGGC	14050
AACCTGGGCT	CCATGGTTGC	ACTAAACGCC	TTCCTGAGTA	CACAGCCCGC	14100
CAACGTGCCG	CGGGGACAGG	AGGACTACAC	CAACTTTGTG	AGCGCACTGC	14150
GGCTAATGGT	GACTGAGACA	CCGCAAAGTG	AGGTGTACCA	GTCTGGGCCA	14200
GACTATTTTT	TCCAGACCAG	TAGACAAGGC	CTGCAGACCG	TAAACCTGAG	14250
CCAGGCTTTC	AAAAACTTGC	AGGGGCTGTG	GGGGGTGCGG	GCTCCCACAG	14300
GCGACCGCGC	GACCGTGTCT	AGCTTGCTGA	CGCCCAACTC	GCGCCTGTTG	14350
CTGCTGCTAA	TAGCGCCCTT	CACGGACAGT	GGCAGCGTGT	CCCGGGACAC	14400
ATACCTAGGT	CACTTGCTGA	CACTGTACCG	CGAGGCCATA	GGTCAGGCGC	14450
ATGTGGACGA	GCATACTTTC	CAGGAGATTA	CAAGTGTGAG	CCGCGCGCTG	14500
GGGCAGGAGG	ACACGGGCAG	CCTGGAGGCA	ACCCTAAACT	ACCTGCTGAC	14550
CAACCGGCGG	CAGAAGATCC	CCTCGTTGCA	CAGTTTAAAC	AGCGAGGAGG	14600

AGCGCATTTT	GCGCTACGTG	CAGCAGAGCG	TGAGCCTTAA	CCTGATGCGC	14650
GACGGGGTAA	CGCCCAGCGT	GGCGCTGGAC	ATGACCGCGC	GCAACATGGA	14700
ACCGGGCATG	TATGCCTCAA	ACCGGCCGTT	TATCAACCGC	CTAATGGACT	14750
ACTTGCATCG	CGCGGCCGCC	GTGAACCCCG	AGTATTTTAC	CAATGCCATC	14800
TTGAACCCGC	ACTGGCTACC	GCCCCCTGGT	TTCTACACCG	GGGGATTCTGA	14850
GGTGCCCGAG	GGTAACGATG	GATTCCTCTG	GGACGACATA	GACGACAGCG	14900
TGTTTTTCCCC	GCAACCGCAG	ACCCTGCTAG	AGTTGCAACA	GCGCGAGCAG	14950
GCAGAGGCGG	CGCTGCGAAA	GGAAAGCTTC	CGCAGGCCAA	GCAGCTTGTC	15000
CGATCTAGGC	GCTGCGGCC	CGCGGTCAGA	TGCTAGTAGC	CCATTTCCAA	15050
GCTTGATAGG	GTCTCTTACC	AGCACTCGCA	CCACCCGCCC	GCGCCTGCTG	15100
GGCGAGGAGG	AGTACCTAAA	CAACTCGCTG	CTGCAGCCGC	AGCGCGAAAA	15150
AAACCTGCCT	CCGGCATTTC	CCAACAACGG	GATAGAGAGC	CTAGTGGACA	15200
AGATGAGTAG	ATGGAAGACG	TACGCGCAGG	AGCACAGGGA	CGTGCCAGGC	15250
CCGCGCCCCG	CCACCCGTCG	TCAAAGGCAC	GACCGTCAGC	GGGGTCTGGT	15300
GTGGGAGGAC	GATGACTCGG	CAGACGACAG	CAGCGTCCTG	GATTTGGGAG	15350
GGAGTGGA	CCCGTTTGCG	CACCTTCGCC	CCAGGCTGGG	GAGAATGTTT	15400
TAAAAA	AAAGCATGAT	GCAAATAAA	AAACTCACCA	AGGCCATGGC	15450
ACCGAGCGTT	GGTTTTCTTG	TATTCCCCTT	AGTATGCGGC	GCGCGGCGAT	15500
GTATGAGGAA	GGTCCTCCTC	CCTCCTACGA	GAGTGTGGTG	AGCGCGGCGC	15550
CAGTGGCGGC	GGCGCTGGGT	TCTCCCTTCG	ATGCTCCCCT	GGACCCGCCG	15600
TTTGTGCCTC	CGCGGTACCT	GCGGCCTACC	GGGGGGAGAA	ACAGCATCCG	15650
TTACTCTGAG	TTGGCACCCC	TATTCGACAC	CACCCGTGTG	TACCTGGTGG	15700
ACAACAAGTC	AACGGATGTG	GCATCCCTGA	ACTACCAGAA	CGACCACAGC	15750
AACTTTCTGA	CCACGGTCAT	TCAAAACAAT	GACTACAGCC	CGGGGGAGGC	15800
AAGCACACAG	ACCATCAATC	TTGACGACCG	GTCGCACTGG	GGCGGCGACC	15850
TGAAAACCAT	CCTGCATACC	AACATGCCAA	ATGTGAACGA	GTTCATGTTT	15900



ACCAATAAGT	TTAAGGCGCG	GGTGATGGTG	TCGCGCTTGC	CTACTAAGGA	15950
CAATCAGGTG	GAGCTGAAAT	ACGAGTGGGT	GGAGTTCACG	CTGCCCCGAGG	16000
GCAACTACTC	CGAGACCATG	ACCATAGACC	TTATGAACAA	CGCGATCGTG	16050
GAGCACTACT	TGAAAGTGGG	CAGACAGAAC	GGGGTTCTGG	AAAGCGACAT	16100
CGGGGTAAAG	TTTGACACCC	GCAACTTCAG	ACTGGGGTTT	GACCCCGTCA	16150
CTGGTCTTGT	CATGCCTGGG	GTATATACAA	ACGAAGCCTT	CCATCCAGAC	16200
ATCATTTTGC	TGCCAGGATG	CGGGGTGGAC	TTCACCCACA	GCCGCCTGAG	16250
CAACTTGTTG	GGCATCCGCA	AGCGGCAACC	CTTCCAGGAG	GGCTTTAGGA	16300
TCACCTACGA	TGATCTGGAG	GGTGGAACA	TTCCCGCACT	GTTGGATGTG	16350
GACGCCTACC	AGGCGAGCTT	GAAAGATGAC	ACCGAACAGG	GCGGGGGTGG	16400
CGCAGGCGGC	AGCAACAGCA	GTGGCAGCGG	CGCGGAAGAG	AACTCCAACG	16450
CGGCAGCCGC	GGCAATGCAG	CCGGTGGAGG	ACATGAACGA	TCATGCCATT	16500
CGCGGCGACA	CCTTTGCCAC	ACGGGCTGAG	GAGAAGCGCG	CTGAGGCCGA	16550
AGCAGCGGCC	GAAGCTGCCG	CCCCCGCTGC	GCAACCCGAG	GTCGAGAAGC	16600
CTCAGAAGAA	ACCGGTGATC	AAACCCCTGA	CAGAGGACAG	CAAGAAACGC	16650
AGTTACAACC	TAATAAGCAA	TGACAGCACC	TTCACCCAGT	ACCGCAGCTG	16700
GTACCTTGCA	TACAACTACG	GCGACCCTCA	GACCGGAATC	CGCTCATGGA	16750
CCCTGCTTTG	CACTCCTGAC	GTAACCTGCG	GCTCGGAGCA	GGTCTACTGG	16800
TCGTTGCCAG	ACATGATGCA	AGACCCCGTG	ACCTTCCGCT	CCACGCGCCA	16850
GATCAGCAAC	TTTCCGGTGG	TGGGCGCCGA	GCTGTTGCCC	GTGCACTCCA	16900
AGAGCTTCTA	CAACGACCAG	GCCGTCTACT	CCCAACTCAT	CCGCCAGTTT	16950
ACCTCTCTGA	CCCACGTGTT	CAATCGCTTT	CCCGAGAACC	AGATTTTGGC	17000
GCGCCCGCCA	GCCCCACCA	TCACCACCGT	CAGTGAAAAC	GTTCTTGCTC	17050
TCACAGATCA	CGGGACGCTA	CCGCTGCGCA	ACAGCATCGG	AGGAGTCCAG	17100
CGAGTGACCA	TTACTGACGC	CAGACGCCGC	ACCTGCCCCCT	ACGTTTACAA	17150
GGCCCTGGGC	ATAGTCTCGC	CGCGCGTCCT	ATCGAGCCGC	ACTTTTTGAG	17200

CAAGCATGTC	CATCCTTATA	TCGCCCAGCA	ATAACACAGG	CTGGGGCCTG	17250
CGCTTCCCAA	GCAAGATGTT	TGGCGGGGCC	AAGAAGCGCT	CCGACCAACA	17300
CCCAGTGCGC	GTGCGCGGGC	ACTACCGCGC	GCCCTGGGGC	GCGCACAAAC	17350
GCGGCCGCAC	TGGGCGCACC	ACCGTCGATG	ACGCCATCGA	CGCGGTGGTG	17400
GAGGAGGCGC	GCAACTACAC	GCCCACGCCG	CCACCAGTGT	CCACAGTGGA	17450
CGCGGCCATT	CAGACCGTGG	TGCGCGGAGC	CCGGCGCTAT	GCTAAAATGA	17500
AGAGACGGCG	GAGGCGCGTA	GCACGTCGCC	ACCGCCGCCG	ACCCGGCACT	17550
GCCGCCCAAC	GCGCGGCGGC	GGCCCTGCTT	AACCGCGCAC	GTCGCACCGG	17600
CCGACGGGCG	GCCATGCGGG	CCGCTCGAAG	GCTGGCCGCG	GGTATTGTCA	17650
CTGTGCCCCC	CAGGTCCAGG	CGACGAGCGG	CCGCCGCAGC	AGCCGCGGCC	17700
ATTAGTGCTA	TGACTCAGGG	TCGCAGGGGC	AACGTGTATT	GGGTGCGCGA	17750
CTCGGTTAGC	GGCCTGCGCG	TGCCCCGTGCG	CACCCGCCCC	CCGCGCAACT	17800
AGATTGCAAG	AAAAAACTAC	TTAGACTCGT	ACTGTTGTAT	GTATCCAGCG	17850
GCGGCGGCGC	GCAACGAAGC	TATGTCCAAG	CGCAAAATCA	AAGAAGAGAT	17900
GCTCCAGGTC	ATCGCGCCGG	AGATCTATGG	CCCCCCGAAG	AAGGAAGAGC	17950
AGGATTACAA	GCCCCGAAAG	CTAAAGCGGG	TCAAAAAGAA	AAAGAAAGAT	18000
GATGATGATG	AACTTGACGA	CGAGGTGGAA	CTGCTGCACG	CTACCGCGCC	18050
CAGGCGACGG	GTACAGTGGA	AAGGTCGACG	CGTAAAACGT	GTTTTGCGAC	18100
CCGGCACCAC	CGTAGTCTTT	ACGCCCGGTG	AGCGCTCCAC	CCGCACCTAC	18150
AAGCGCGTGT	ATGATGAGGT	GTACGGCGAC	GAGGACCTGC	TTGAGCAGGC	18200
CAACGAGCGC	CTCGGGGAGT	TTGCCTACGG	AAAGCGGCAT	AAGGACATGC	18250
TGGCGTTGCC	GCTGGACGAG	GGCAACCCAA	CACCTAGCCT	AAAGCCCGTA	18300
ACACTGCAGC	AGGTGCTGCC	CGCGCTTGCA	CCGTCCGAAG	AAAAGCGCGG	18350
CCTAAAGCGC	GAGTCTGGTG	ACTTGGCACC	CACCGTGCAG	CTGATGGTAC	18400
CCAAGCGCCA	GCGACTGGAA	GATGTCTTGG	AAAAAATGAC	CGTGGAACCT	18450
GGGCTGGAGC	CCGAGGTCCG	CGTGCGGCCA	ATCAAGCAGG	TGGCGCCGGG	18500

ACTGGGCGTG	CAGACCGTGG	ACGTTTCAGAT	ACCCACTACC	AGTAGCACCA	18550
GTATTGCCAC	CGCCACAGAG	GGCATGGAGA	CACAAACGTC	CCCGGTTGCC	18600
TCAGCGGTGG	CGGATGCCGC	GGTGCAGGCG	GTCGCTGCGG	CCGCGTCCAA	18650
GACCTCTACG	GAGGTGCAAA	CGGACCCGTG	GATGTTTCGC	GTTTCAGCCC	18700
CCCGGCGCCC	GCGCGGTTCG	AGGAAGTACG	GCGCCGCCAG	CGCGCTACTG	18750
CCCGAATATG	CCCTACATCC	TTCCATTGCG	CCTACCCCCG	GCTATCGTGG	18800
CTACACCTAC	CGCCCCAGAA	GACGAGCAAC	TACCCGACGC	CGAACCACCA	18850
CTGGAACCCG	CCGCCGCCGT	CGCCGTGCGC	AGCCCGTGCT	GGCCCCGATT	18900
TCCGTGCGCA	GGGTGGCTCG	CGAAGGAGGC	AGGACCCTGG	TGCTGCCAAC	18950
AGCGCGCTAC	CACCCCAGCA	TCGTTTAAAA	GCCGGTCTTT	GTGGTTCTTG	19000
CAGATATGGC	CCTCACCTGC	CGCCTCCGTT	TCCCGGTGCC	GGGATTCCGA	19050
GGAAGAATGC	ACCGTAGGAG	GGGCATGGCC	GGCCACGGCC	TGACGGGCGG	19100
CATGCGTCGT	GCGCACCACC	GGCGGCGGCG	CGCGTCGCAC	CGTCGCATGC	19150
GCGGCGGTAT	CCTGCCCCCTC	CTTATTCCAC	TGATCGCCGC	GGCGATTGGC	19200
GCCGTGCCCG	GAATTGCATC	CGTGGCCTTG	CAGGCGCAGA	GACACTGATT	19250
AAAAACAAGT	TGCATGTGGA	AAAATCAAAA	TAAAAAGTCT	GGACTCTCAC	19300
GCTCGCTTGG	TCCTGTAACT	ATTTTGTAGA	ATGGAAGACA	TCAACTTTGC	19350
GTCTCTGGCC	CCGCGACACG	GCTCGCGCCC	GTTTCATGGGA	AACTGGCAAG	19400
ATATCGGCAC	CAGCAATATG	AGCGGTGGCG	CCTTCAGCTG	GGGCTCGCTG	19450
TGGAGCGGCA	TTAAAAATTT	CGGTTCCACC	GTTAAGAACT	ATGGCAGCAA	19500
GGCCTGGAAC	AGCAGCACAG	GCCAGATGCT	GAGGGATAAG	TTGAAAGAGC	19550
AAAATTTCCA	ACAAAAGGTG	GTAGATGGCC	TGGCCTCTGG	CATTAGCGGG	19600
GTGGTGGACC	TGGCCAACCA	GGCAGTGCAA	AATAAGATTA	ACAGTAAGCT	19650
TGATCCCCGC	CCTCCCGTAG	AGGAGCCTCC	ACCGGCCGTG	GAGACAGTGT	19700
CTCCAGAGGG	GCGTGGCGAA	AAGCGTCCGC	GCCCCGACAG	GGAAGAACT	19750
CTGGTGACGC	AAATAGACGA	GCCTCCCTCG	TACGAGGAGG	CACTAAAGCA	19800

AGGCCTGCCC	ACCACCCGTC	CCATCGCGCC	CATGGCTACC	GGAGTGCTGG	19850
GCCAGCACAC	ACCCGTAACG	CTGGACCTGC	CTCCCCCGC	CGACACCCAG	19900
CAGAAACCTG	TGCTGCCAGG	CCCGACCGCC	GTTGTTGTAA	CCCGTCCTAG	19950
CCGCGCGTCC	CTGCGCCGCG	CCGCCAGCGG	TCCGCGATCG	TTGCGGCCCC	20000
TAGCCAGTGG	CAACTGGCAA	AGCACACTGA	ACAGCATCGT	GGGTCTGGGG	20050
GTGCAATCCC	TGAAGCGCCG	ACGATGCTTC	TGAATAGCTA	ACGTGTCGTA	20100
TGTGTGTCAT	GTATGCGTCC	ATGTGCGCCG	CAGAGGAGCT	GCTGAGCCGC	20150
CGCGCGCCCC	CTTTCCAAGA	TGGCTACCCC	TTCGATGATG	CCGCAGTGGT	20200
CTTACATGCA	CATCTCGGGC	CAGGACGCCT	CGGAGTACCT	GAGCCCCGGG	20250
CTGGTGCACT	TTGCCCCGCG	CACCGAGACG	TACTTCAGCC	TGAATAACAA	20300
GTTTAGAAAC	CCCACGGTGG	CGCCTACGCA	CGACGTGACC	ACAGACCGGT	20350
CCCAGCGTTT	GACGCTGCGG	TTCATCCCTG	TGGACCGTGA	GGATACTGCG	20400
TACTCGTACA	AGGCGCGGTT	CACCCTAGCT	GTGGGTGATA	ACCGTGTGCT	20450
GGACATGGCT	TCCACGTACT	TTGACATCCG	CGGCGTGCTG	GACAGGGGCC	20500
CTACTTTTAA	GCCCTACTCT	GGCACTGCCT	ACAACGCCCT	GGCTCCCAAG	20550
GGTGCCCCAA	ATCCTTGCGA	ATGGGATGAA	GCTGCTACTG	CTCTTGAAAT	20600
AAACCTAGAA	GAAGAGGACG	ATGACAACGA	AGACGAAGTA	GACGAGCAAG	20650
CTGAGCAGCA	AAAAACTCAC	GTATTTGGGC	AGGCGCCTTA	TTCTGGTATA	20700
AATATTACAA	AGGAGGGTAT	TCAAATAGGT	GTCGAAGGTC	AAACACCTAA	20750
ATATGCCGAT	AAAACATTTT	AACCTGAACC	TCAAATAGGA	GAATCTCAGT	20800
GGTACGAAAC	TGAAATTAAT	CATGCAGCTG	GGAGAGTCCT	TAAAAAGACT	20850
ACCCCAATGA	AACCATGTTA	CGGTTTATAT	GCAAAACCCA	CAAATGAAAA	20900
TGGAGGGCAA	GGCATTCTTG	TAAAGCAACA	AAATGGAAAG	CTAGAAAGTC	20950
AAGTGGAAT	GCAATTTTTC	TCAACTACTG	AGGCGACCGC	AGGCAATGGT	21000
GATAACTTGA	CTCCTAAAGT	GGTATTGTAC	AGTGAAGATG	TAGATATAGA	21050
AACCCAGAC	ACTCATATTT	CTTACATGCC	CACTATTAAG	GAAGGTAAGT	21100

CACGAGAACT AATGGGCCAA CAATCTATGC CCAACAGGCC TAATTACATT 21150  
GCTTTTAGGG ACAATTTTAT TGGTCTAATG TATTACAACA GCACGGGTAA 21200  
TATGGGTGTT CTGGCGGGCC AAGCATCGCA GTTGAATGCT GTTGTAGATT 21250  
TGCAAGACAG AAACACAGAG CTTTCATACC AGCTTTTGCT TGATTCCATT 21300  
GGTGATAGAA CCAGGTACTT TTCTATGTGG AATCAGGCTG TTGACAGCTA 21350  
TGATCCAGAT GTTAGAATTA TTGAAAATCA TGGAACTGAA GATGAACTTC 21400  
CAAATTACTG CTTTCCACTG GGAGGTGTGA TTAATACAGA GACTCTTACC 21450  
AAGGTAAAAC CTAAAACAGG TCAGGAAAAT GGATGGGAAA AAGATGCTAC 21500  
AGAATTTTCA GATAAAAATG AAATAAGAGT TGGAAATAAT TTTGCCATGG 21550  
AAATCAATCT AAATGCCAAC CTGTGGAGAA ATTTCTGTGA CTCCAACATA 21600  
GCGCTGTATT TGCCCGACAA GCTAAAGTAC AGTCCTTCCA ACGTAAAAAT 21650  
TTCTGATAAC CCAAACACCT ACGACTACAT GAACAAGCGA GTGGTGGCTC 21700  
CCGGGTAGT GGACTGCTAC ATTAACCTTG GAGCACGCTG GTCCCTTGAC 21750  
TATATGGACA ACGTCAACCC ATTTAACCAC CACCGCAATG CTGGCCTGCG 21800  
CTACCGCTCA ATGTTGCTGG GCAATGGTCG CTATGTGCCC TTCCACATCC 21850  
AGGTGCCTCA GAAGTTCTTT GCCATTAAAA ACCTCCTTCT CCTGCCGGGC 21900  
TCATACACCT ACGAGTGGAA CTTAGGAAG GATGTTAACA TGGTTCTGCA 21950  
GAGCTCCCTA GGAAATGACC TAAGGGTTGA CGGAGCCAGC ATTAAGTTTG 22000  
ATAGCATTTG CCTTTACGCC ACCTTCTTCC CCATGGCCCA CAACACCGCC 22050  
TCCACGCTTG AGGCCATGCT TAGAAACGAC ACCAACGACC AGTCCTTTAA 22100  
CGACTATCTC TCCGCCGCCA ACATGCTCTA CCCTATACCC GCCAACGCTA 22150  
CCAACGTGCC CATATCCATC CCCTCCCGCA ACTGGGCGGC TTTCCGCGGC 22200  
TGGGCCTTCA CGCGCCTTAA GACTAAGGAA ACCCCATCAC TGGGCTCGGG 22250  
CTACGACCCT TATTACACCT ACTCTGGCTC TATACCCTAC CTAGATGGAA 22300  
CCTTTTACCT CAACCACACC TTTAAGAAGG TGGCCATTAC CTTTGA CTCT 22350  
TCTGTCAGCT GGCCTGGCAA TGACCGCCTG CTTACCCCA ACGAGTTTGA 22400

AATTAAGCGC TCAGTTGACG GGGAGGGTTA CAACGTTGCC CAGTGTAACA 22450  
TGACCAAAGA CTGGTTCCTG GTACAAATGC TAGCTAACTA CAACATTGGC 22500  
TACCAGGGCT TCTATATCCC AGAGAGCTAC AAGGACCGCA TGTACTCCTT 22550  
CTTTAGAAAC TTCCAGCCCA TGAGCCGTC A GGTGGTGGAT GATACTAAAT 22600  
ACAAGGACTA CCAACAGGTG GGCATCCTAC ACCAACACAA CAACTCTGGA 22650  
TTTGTGGCT ACCTTGCCCC CACCATGCGC GAAGGACAGG CCTACCCTGC 22700  
TAACTTCCCC TATCCGCTTA TAGGCAAGAC CGCAGTTGAC AGCATTACCC 22750  
AGAAAAAGTT TCTTTGCGAT CGCACCTTT GCGCATCCC ATTCTCCAGT 22800  
AACTTTATGT CCATGGGCGC ACTCACAGAC CTGGGCCAAA ACCTTCTCTA 22850  
CGCCAACTCC GCCCAGCGC TAGACATGAC TTTTGAGGTG GATCCCATGG 22900  
ACGAGCCCAC CCTTCTTTAT GTTTTGTTTG AAGTCTTTGA CGTGGTCCGT 22950  
GTGCACCGGC CGCACCGCG CGTCATCGAA ACCGTGTACC TGCGCACGCC 23000  
CTTCTCGGCC GGCAACGCCA CAACATAAAG AAGCAAGCAA CATCAACAAC 23050  
AGCTGCCGCC ATGGGCTCCA GTGAGCAGGA ACTGAAAGCC ATTGTCAAAG 23100  
ATCTTGGTTG TGGGCCATAT TTTTGGGCA CCTATGACAA GCGCTTTCCA 23150  
GGCTTTGTTT CTCCACACAA GCTCGCCTGC GCCATAGTCA ATACGGCCGG 23200  
TCGCGAGACT GGGGGCGTAC ACTGGATGGC CTTTGCCTGG AACCCGCACT 23250  
CAAAAACATG CTACCTCTTT GAGCCCTTTG GCTTTTCTGA CCAGCGACTC 23300  
AAGCAGGTTT ACCAGTTTGA GTACGAGTCA CTCCTGCGCC GTAGCGCCAT 23350  
TGCTTCTTCC CCCGACCGCT GTATAACGCT GGAAAAGTCC ACCCAAAGCG 23400  
TACAGGGGCC CAACTCGGCC GCCTGTGGAC TATTCTGCTG CATGTTTCTC 23450  
CACGCCTTTG CCAACTGGCC CCAAACTCCC ATGGATCACA ACCCCACCAT 23500  
GAACCTTATT ACCGGGGTAC CCAACTCCAT GCTCAACAGT CCCCAGGTAC 23550  
AGCCCACCCT GCGTCGCAAC CAGGAACAGC TCTACAGCTT CCTGGAGCGC 23600  
CACTCGCCCT ACTTCCGCAG CCACAGTGCG CAGATTAGGA GCGCCACTTC 23650  
TTTTTGTCAC TTGAAAAACA TGTA AAAATA ATGTACTAGA GAACTTTCA 23700

ATAAAGGCAA	ATGCTTTTAT	TTGTACACTC	TCGGGTGATT	ATTTACCCCC	23750
ACCCCTGCCG	TCTGCGCCGT	TTAAAAATCA	AAGGGGTTC	GCCGCGCATC	23800
GCTATGCGCC	ACTGGCAGGG	ACACGTTGCG	ATACTGGTGT	TTAGTGCTCC	23850
ACTTAAACTC	AGGCACAACC	ATCCGCGGCA	GCTCGGTGAA	GTTTTCACTC	23900
CACAGGCTGC	GCACCATCAC	CAACGCGTTT	AGCAGGTCGG	GCGCCGATAT	23950
CTTGAAGTCG	CAGTTGGGGC	CTCCGCCCTG	CGCGCGCGAG	TTGCGATACA	24000
CAGGGTTGCA	GCACTGGAAC	ACTATCAGCG	CCGGGTGGTG	CACGCTGGCC	24050
AGCACGCTCT	TGTCGGAGAT	CAGATCCGCG	TCCAGGTCCT	CCGCGTTGCT	24100
CAGGGCGAAC	GGAGTCAACT	TTGGTAGCTG	CCTTCCCAA	AAGGGCGCGT	24150
GCCCAGGCTT	TGAGTTGCAC	TCGCACCGTA	GTGGCATCAA	AAGGTGACCG	24200
TGCCCCGGTCT	GGGCGTTAGG	ATACAGCGCC	TGCATAAAAG	CCTTGATCTG	24250
CTTAAAAGCC	ACCTGAGCCT	TTGCGCCTTC	AGAGAAGAAC	ATGCCGCAAG	24300
ACTTGCCGGA	AAACTGATTG	GCCGGACAGG	CCGCGTCGTG	CACGCAGCAC	24350
CTTGCGTCGG	TGTTGGAGAT	CTGCACCACA	TTTCGGCCCC	ACCGTTCTTT	24400
CACGATCTTG	GCCTTGCTAG	ACTGCTCCTT	CAGCGCGCGC	TGCCCCTTTT	24450
CGCTCGTCAC	ATCCATTTCA	ATCACGTGCT	CCTTATTTAT	CATAATGCTT	24500
CCGTGTAGAC	ACTTAAGCTC	GCCTTCGATC	TCAGCGCAGC	GGTGCAGCCA	24550
CAACGCGCAG	CCCGTGGGCT	CGTGATGCTT	GTAGGTCACC	TCTGCAAACG	24600
ACTGCAGGTA	CGCCTGCAGG	AATCGCCCCA	TCATCGTCAC	AAAGGTCTTG	24650
TTGCTGGTGA	AGGTCAGCTG	CAACCCGCGG	TGCTCCTCGT	TCAGCCAGGT	24700
CTTGCATACG	GCCGCCAGAG	CTTCCACTTG	GTCAGGCAGT	AGTTTGAAGT	24750
TCGCCTTTAG	ATCGTTATCC	ACGTGGTACT	TGTCCATCAG	CGCGCGCGCA	24800
GCCTCCATGC	CCTTCTCCCA	CGCAGACACG	ATCGGCACAC	TCAGCGGGTT	24850
CATCACCGTA	ATTTCACTTT	CCGCTTCGCT	GGGCTCTTCC	TCTTCCTCTT	24900
GCGTCCGCAT	ACCACGCGCC	ACTGGGTGCT	CTTCATTCAG	CCGCCGCACT	24950
GTGCGCTTAC	CTCCTTTGCC	ATGCTTGATT	AGCACCGGTG	GGTTGCTGAA	25000

ACCCACCATT	TGTAGCGCCA	CATCTTCTCT	TTCTTCCTCG	CTGTCCACGA	25050
TTACCTCTGG	TGATGGCGGG	CGCTCGGGCT	TGGGAGAAGG	GCGCTTCTTT	25100
TTCTTCTTGG	GCGCAATGGC	CAAATCCGCC	GCCGAGGTCG	ATGGCCGCGG	25150
GCTGGGTGTG	CGCGGCACCA	GCGCGTCTTG	TGATGAGTCT	TCCTCGTCCT	25200
CGGACTCGAT	ACGCCGCCTC	ATCCGCTTTT	TTGGGGGCGC	CCGGGGAGGC	25250
GGCGGCGACG	GGGACGGGGA	CGACACGTCC	TCCATGGTTG	GGGGACGTGC	25300
CGCCGCACCG	CGTCCGCGCT	CGGGGGTGGT	TTCGCGCTGC	TCCTCTTCCC	25350
GACTGGCCAT	TTCCTTCTCC	TATAGGCAGA	AAAAGATCAT	GGAGTCAGTC	25400
GAGAAGAAGG	ACAGCCTAAC	CGCCCCCTCT	GAGTTCGCCA	CCACCGCCTC	25450
CACCGATGCC	GCCAACGCGC	CTACCACCTT	CCCCGTCGAG	GCACCCCCGC	25500
TTGAGGAGGA	GGAAGTGATT	ATCGAGCAGG	ACCCAGGTTT	TGTAAGCGAA	25550
GACGACGAGG	ACCGCTCAGT	ACCAACAGAG	GATAAAAAGC	AAGACCAGGA	25600
CAACGCAGAG	GCAAACGAGG	AACAAGTCGG	GCGGGGGGAC	GAAAGGCATG	25650
GCGACTACCT	AGATGTGGGA	GACGACGTGC	TGTTGAAGCA	TCTGCAGCGC	25700
CAGTGCGCCA	TTATCTGCGA	CGCGTTGCAA	GAGCGCAGCG	ATGTGCCCCT	25750
CGCCATAGCG	GATGTCAGCC	TTGCCTACGA	ACGCCACCTA	TTCTCACCGC	25800
GCGTACCCCC	CAAACGCCAA	GAAAACGGCA	CATGCGAGCC	CAACCCGCGC	25850
CTCAACTTCT	ACCCCGTATT	TGCCGTGCCA	GAGGTGCTTG	CCACCTATCA	25900
CATCTTTTTT	CAAAACTGCA	AGATACCCCT	ATCCTGCCGT	GCCAACCGCA	25950
GCCGAGCGGA	CAAGCAGCTG	GCCTTGCGGC	AGGGCGCTGT	CATACCTGAT	26000
ATCGCCTCGC	TCAACGAAGT	GCCAAAAATC	TTTGAGGGTC	TTGGACGCGA	26050
CGAGAAGCGC	GCGGCAAACG	CTCTGCAACA	GGAAAACAGC	GAAAATGAAA	26100
GTCACCTCTG	AGTGTTGGTG	GAAGTCGAGG	GTGACAACGC	GCGCCTAGCC	26150
GTAATAAAAC	GCAGCATCGA	GGTCACCCAC	TTTGCCTACC	CGGCACTTAA	26200
CCTACCCCCC	AAGGTCATGA	GCACAGTCAT	GAGTGAGCTG	ATCGTGCGCC	26250
GTGCGCAGCC	CCTGGAGAGG	GATGCAAATT	TGCAAGAACA	AACAGAGGAG	26300



GGCCTACCCG CAGTTGGCGA CGAGCAGCTA GCGCGCTGGC TTCAAACGCG 26350  
CGAGCCTGCC GACTTGGAGG AGCGACGCAA ACTAATGATG GCCGCAGTGC 26400  
TCGTTACCGT GGAGCTTGAG TGCATGCAGC GGTTCCTTTC TGACCCGGAG 26450  
ATGCAGCGCA AGCTAGAGGA AACATTGCAC TACACCTTTC GACAGGGCTA 26500  
CGTACGCCAG GCCTGCAAGA TCTCCAACGT GGAGCTCTGC AACCTGGTCT 26550  
CCTACCTTGG AATTTTGCAC GAAAACCGCC TTGGGCAAAA CGTGCTTCAT 26600  
TCCACGCTCA AGGGCGAGGC GCGCCGCGAC TACGTCCGCG ACTGCGTTTA 26650  
CTTATTTCTA TGCTACACCT GGCAGACGGC CATGGGCGTT TGGCAGCAGT 26700  
GCTTGGAGGA GTGCAACCTC AAGGAGCTGC AGAACTGCT AAAGCAAAAC 26750  
TTGAAGGACC TATGGACGGC CTTCAACGAG CGCTCCGTGG CCGCGCACCT 26800  
GGCGGACATC ATTTTCCCCG AACGCCTGCT TAAAACCCTG CAACAGGGTC 26850  
TGCCAGACTT CACCAGTCAA AGCATGTTGC AGAACTTTAG GAACTTTATC 26900  
CTAGAGCGCT CAGGAATCTT GCGCGCCACC TGCTGTGCAC TTCCTAGCGA 26950  
CTTTGTGCCC ATTAAGTACC GCGAATGCCC TCCGCCGCTT TGGGGCCACT 27000  
GCTACCTTCT GCAGCTAGCC AACTACCTTG CCTACCACTC TGACATAATG 27050  
GAAGACGTGA GCGGTGACGG TCTACTGGAG TGTCAGTGC GCTGCAACCT 27100  
ATGCACCCCG CACCGCTCCC TGGTTTGCAA TTCGCAGCTG CTTAACGAAA 27150  
GTCAAATTAT CGGTACCTTT GAGCTGCAGG GTCCCTCGCC TGACGAAAAG 27200  
TCCGCGGCTC CGGGGTGAA ACTCACTCCG GGGCTGTGGA CGTCGGCTTA 27250  
CCTTCGCAAA TTTGTACCTG AGGACTACCA CGCCCACGAG ATTAGGTTCT 27300  
ACGAAGACCA ATCCCGCCCG CCAAATGCGG AGCTTACCGC CTGCGTCATT 27350  
ACCCAGGGCC ACATTCTTGG CCAATTGCAA GCCATCAACA AAGCCCGCCA 27400  
AGAGTTTCTG CTACGAAAGG GACGGGGGGT TTAATTGGAC CCCCAGTCCG 27450  
GCGAGGAGCT CAACCCAATC CCCCCGCCG CGCAGCCCTA TCAGCAGCAG 27500  
CCGCGGGCCC TTGCTTCCCA GGATGGCACC CAAAAGAAG CTGCAGCTGC 27550  
CGCCGCCACC CACGGACGAG GAGGAATACT GGGACAGTCA GGCAGAGGAG 27600

GTTTTGGACG	AGGAGGAGGA	GGACATGATG	GAAGACTGGG	AGAGCCTAGA	27650
CGAGGAAGCT	TCCGAGGTCG	AAGAGGTGTC	AGACGAAACA	CCGTCACCCT	27700
CGGTTCGCATT	CCCCTCGCCG	GCGCCCCAGA	AATCGGCAAC	CGGTTCCAGC	27750
ATGGCTACAA	CCTCCGCTCC	TCAGGCGCCG	CCGGCACTGC	CCGTTGCGCCG	27800
ACCCAACCGT	AGATGGGACA	CCACTGGAAC	CAGGGCCGGT	AAGTCCAAGC	27850
AGCCGCGGCC	GTTAGCCCAA	GAGCAACAAC	AGCGCCAAGG	CTACCGCTCA	27900
TGGCGCGGGC	ACAAGAACGC	CATAGTTGCT	TGCTTGCAAG	ACTGTGGGGG	27950
CAACATCTCC	TTCGCCCCGCC	GCTTTCTTCT	CTACCATCAC	GGCGTGGCCT	28000
TCCCCCGTAA	CATCCTGCAT	TACTACCGTC	ATCTCTACAG	CCCATACTGC	28050
ACCGGCGGCA	GCGGCAGCGG	CAGCAACAGC	AGCGGCCACA	CAGAAGCAAA	28100
GGCGACCGGA	TAGCAAGACT	CTGACAAAGC	CCAAGAAATC	CACAGCGGCG	28150
GCAGCAGCAG	GAGGAGGAGC	GCTGCGTCTG	GCGCCCAACG	AACCCGTATC	28200
GACCCGCGAG	CTTAGAAACA	GGATTTTTTCC	CACTCTGTAT	GCTATATTTT	28250
AACAGAGCAG	GGGCCAAGAA	CAAGAGCTGA	AAATAAAAAA	CAGGTCTCTG	28300
CGATCCCTCA	CCCGCAGCTG	CCTGTATCAC	AAAAGCGAAG	ATCAGCTTCG	28350
GCGCACGCTG	GAAGACGCGG	AGGCTCTCTT	CAGTAAATAC	TGCGCGCTGA	28400
CTCTTAAGGA	CTAGTTTCGC	GCCCTTTCTC	AAATTTAAGC	GCGAAACTA	28450
CGTCATCTCC	AGCGGCCACA	CCCGGCGCCA	GCACCTGTCT	TCAGCGCCAT	28500
TATGAGCAAG	GAAATTCCCA	CGCCCTACAT	GTGGAGTTAC	CAGCCACAAA	28550
TGGGACTTGC	GGCTGGAGCT	GCCCAAGACT	ACTCAACCCG	AATAAACTAC	28600
ATGAGCGCGG	GACCCACAT	GATATCCCGG	GTCAACGGAA	TCCGCGCCCA	28650
CCGAAACCGA	ATTCTCTTGG	AACAGGCGGC	TATTACCACC	ACACCTCGTA	28700
ATAACCTTAA	TCCCCGTAGT	TGGCCCGCTG	CCCTGGTGTA	CCAGGAAAGT	28750
CCCCTCCCA	CCACTGTGGT	ACTTCCCAGA	GACGCCCAGG	CCGAAGTTCA	28800
GATGACTAAC	TCAGGGGCGC	AGCTTGCGGG	CGGCTTTCGT	CACAGGGTGC	28850
GGTCGCCCCG	GCAGGGTATA	ACTCACCTGA	CAATCAGAGG	GCGAGGTATT	28900

CAGCTCAACG ACGAGTCGGT GAGCTCCTCG CTTGGTCTCC GTCCGGACGG 28950  
GACATTTTCAG ATCGGCGGCG CCGGCCGTCC TTCATTACAG CCTCGTCAGG 29000  
CAATCCTAAC TCTGCAGACC TCGTCCTCTG AGCCGCGCTC TGGAGGCATT 29050  
GGAACCTCTGC AATTTATTGA GGAGTTTGTG CCATCGGTCT ACTTTAACCC 29100  
CTTCTCGGGA CCTCCCGGCC ACTATCCGGA TCAATTTATT CCTAACTTTG 29150  
ACGCGGTAAA GGACTCGGCG GACGGCTACG ACTGAATGTT AAGTGGAGAG 29200  
GCAGAGCAAC TGCGCCTGAA ACACCTGGTC CACTGTCGCC GCCACAAGTG 29250  
CTTTGCCCCG GACTCCGGTG AGTTTGTGCTA CTTTGAATTG CCCGAGGATC 29300  
ATATCGAGGG CCCGGCGCAC GCGGTCCGGC TTACCGCCCA GGGAGAGCTT 29350  
GCCCCGTAGCC TGATTCGGGA GTTTACCCAG CGCCCCCTGC TAGTTGAGCG 29400  
GGACAGGGGA CCCTGTGTTC TCACTGTGAT TTGCAACTGT CCTAACCTTG 29450  
GATTACATCA AGATCTTTGT TGCCATCTCT GTGCTGAGTA TAATAAATAC 29500  
AGAAATTAAA ATATACTGGG GCTCCTATCG CCATCCTGTA AACGCCACCG 29550  
TCTTCACCCG CCCAAGCAAA CCAAGGCGAA CCTTACCTGG TACTTTTAAC 29600  
ATCTCTCCCT CTGTGATTTA CAACAGTTTC AACCCAGACG GAGTGAGTCT 29650  
ACGAGAGAAC CTCTCCGAGC TCAGCTACTC CATCAGAAAA AACACCACCC 29700  
TCCTTACCTG CCGGGAACGT ACGAGTGCGT CACCGGCCGC TGCACCACAC 29750  
CTACCGCCTG ACCGTAAACC AGACTTTTTTC CGGACAGACC TCAATAACTC 29800  
TGTTTACCAG AACAGGAGGT GAGCTTAGAA AACCCTTAGG GTATTAGGCC 29850  
AAAGGCGCAG CTA CTGTGGG GTTTATGAAC AATTCAAGCA ACTCTACGGG 29900  
CTATTCTAAT TCAGGTTTCT CTAGAATCGG GGTGTTGGGTT ATTCTCTGTC 29950  
TTGTGATTCT CTTTATTCTT ATACTAACGC TTCTCTGCCT AAGGCTCGCC 30000  
GCCTGCTGTG TGCACATTTG CATTTATTGT CAGCTTTTTTA AACGCTGGGG 30050  
TCGCCACCCA AGATGATTAG GTACATAATC CTAGGTTTAC TCACCCTTGC 30100  
GTCAGCCCAC GGTACCACCC AAAAGGTGGA TTTTAAGGAG CCAGCCTGTA 30150  
ATGTTACATT CGCAGCTGAA GCTAATGAGT GCACCACTCT TATAAAATGC 30200

ACCACAGAAC ATGAAAAGCT GCTTATTCGC CACAAAAACA AAATTGGCAA 30250  
GTATGCTGTT TATGCTATTT GGCAGCCAGG TGACACTACA GAGTATAATG 30300  
TTACAGTTTT CCAGGGTAAA AGTCATAAAA CTTTTATGTA TACTTTTCCA 30350  
TTTTATGAAA TGTGCGACAT TACCATGTAC ATGAGCAAAC AGTATAAGTT 30400  
GTGGCCCCCA CAAAATTGTG TGGAAAACAC TGGCACTTTC TGCTGCACTG 30450  
CTATGCTAAT TACAGTGCTC GCTTTGGTCT GTACCCTACT CTATATTAAA 30500  
TACAAAAGCA GACGCAGCTT TATTGAGGAA AAGAAAATGC CTTAATTTAC 30550  
TAAGTTACAA AGCTAATGTC ACCACTAACT GCTTTACTCG CTGCTTGCAA 30600  
AACAAATTCA AAAAGTTAGC ATTATAATTA GAATAGGATT TAAACCCCCC 30650  
GGTCATTTCC TGCTCAATAC CATTCCCCTG AACAAATTGAC TCTATGTGGG 30700  
ATATGCTCCA GCGCTACAAC CTTGAAGTCA GGCTTCCTGG ATGTCAGCAT 30750  
CTGACTTTGG CCAGCACCTG TCCCGCGGAT TTGTTCCAGT CCAACTACAG 30800  
CGACCCACCC TAACAGAGAT GACCAACACA ACCAACGCGG CCGCCGCTAC 30850  
CGGACTTACA TCTACCACAA ATACACCCCA AGTTTCTGCC TTTGTCAATA 30900  
ACTGGGATAA CTTGGGCATG TGGTGGTTCT CCATAGCGCT TATGTTTGTA 30950  
TGCCTTATTA TTATGTGGCT CATCTGCTGC CTAAAGCGCA AACGCGCCCCG 31000  
ACCACCCATC TATAGTCCCA TCATTGTGCT ACACCCAAAC AATGATGGAA 31050  
TCCATAGATT GGACGGACTG AAACACATGT TCTTTTCTCT TACAGTATGA 31100  
TTAAATGAGA CATGATTCCT CGAGTTTTTA TATTACTGAC CCTTGTTGCG 31150  
CTTTTTTG TG CGTGCTCCAC ATTGGCTGCG GTTTCTCACA TCGAAGTAGA 31200  
CTGCATTCCA GCCTTCACAG TCTATTTGCT TTACGGATTT GTCACCCTCA 31250  
CGCTCATCTG CAGCCTCATC ACTGTGGTCA TCGCCTTTAT CCAGTGCATT 31300  
GACTGGGTCT GTGTGCGCTT TGCATATCTC AGACACCATC CCCAGTACAG 31350  
GGACAGGACT ATAGCTGAGC TTCTTAGAAT TCTTTAATTA TGAAATTTAC 31400  
TGTGACTTTT CTGCTGATTA TTTGCACCCT ATCTGCGTTT TGTTCCCCGA 31450  
CCTCCAAGCC TCAAAGACAT ATATCATGCA GATTCACCTG TATATGGAAT 31500

ATTCCAAGTT	GCTACAATGA	AAAAAGCGAT	CTTTCCGAAG	CCTGGTTATA	31550
TGCAATCATC	TCTGTTATGG	TGTTCTGCAG	TACCATCTTA	GCCCTAGCTA	31600
TATATCCCTA	CCTTGACATT	GGCTGGAAAC	GAATAGATGC	CATGAACCAC	31650
CCAACTTTCC	CCGCGCCCGC	TATGCTTCCA	CTGCAACAAG	TTGTTGCCGG	31700
CGGCTTTGTC	CCAGCCAATC	AGCCTCGCCC	CACCTTCTCCC	ACCCCCACTG	31750
AAATCAGCTA	CTTTAATCTA	ACAGGAGGAG	ATGACTGACA	CCCTAGATCT	31800
AGAAATGGAC	GGAATTATTA	CAGAGCAGCG	CCTGCTAGAA	AGACGCAGGG	31850
CAGCGGCCGA	GCAACAGCGC	ATGAATCAAG	AGCTCCAAGA	CATGGTTAAC	31900
TTGCACCAGT	GCAAAAGGGG	TATCTTTTGT	CTGGTAAAGC	AGGCCAAAGT	31950
CACCTACGAC	AGTAATACCA	CCGGACACCG	CCTTAGCTAC	AAGTTGCCAA	32000
CCAAGCGTCA	GAAATTGGTG	GTCATGGTGG	GAGAAAAGCC	CATTACCATA	32050
ACTCAGCACT	CGGTAGAAAC	CGAAGGCTGC	ATTCACCTCAC	CTTGTCAAGG	32100
ACCTGAGGAT	CTCTGCACCC	TTATTAAGAC	CCTGTGCGGT	CTCAAAGATC	32150
TTATTCCCTT	TAACTAATAA	AAAAAAATAA	TAAAGCATCA	CTTACTTAAA	32200
ATCAGTTAGC	AAATTTCTGT	CCAGTTTATT	CAGCAGCACC	TCCTTGCCCT	32250
CCTCCCAGCT	CTGGTATTGC	AGCTTCCTCC	TGGCTGCAAA	CTTTCTCCAC	32300
AATCTAAATG	GAATGTCAGT	TTCCTCCTGT	TCCTGTCCAT	CCGCACCCAC	32350
TATCTTCATG	TTGTTGCAGA	TGAAGCGCGC	AAGACCGTCT	GAAGATACCT	32400
TCAACCCCGT	GTATCCATAT	GACACGGAAA	CCGGTCCTCC	AACTGTGCCT	32450
TTTCTTACTC	CTCCCTTTGT	ATCCCCCAAT	GGGTTTCAAG	AGAGTCCCCC	32500
TGGGGTACTC	TCTTTGCGCC	TATCCGAACC	TCTAGTTACC	TCCAATGGCA	32550
TGCTTGCGCT	CAAAATGGGC	AACGGCCTCT	CTCTGGACGA	GGCCGGCAAC	32600
CTTACCTCCC	AAAATGTAAC	CACCTGTGAGC	CCACCTCTCA	AAAAAACCAA	32650
GTCAAACATA	AACCTGGAAA	TATCTGCACC	CCTCACAGTT	ACCTCAGAAG	32700
CCCTAACTGT	GGCTGCCGCC	GCACCTCTAA	TGGTCGCGGG	CAACACACTC	32750
ACCATGCAAT	CACAGGCCCC	GCTAACCGTG	CACGACTCCA	AACTTAGCAT	32800

TGCCACCCAA	GGACCCCTCA	CAGTGTGAGA	AGGAAAGCTA	GCCCTGCAAA	32850
CATCAGGCCC	CCTCACCACC	ACCGATAGCA	GTACCCTTAC	TATCACTGCC	32900
TCACCCCCTC	TAATACTGTC	CACTGGTAGC	TTGGGCATTG	ACTTGAAAGA	32950
GCCCATTAT	ACACAAAATG	GAAAACTAGG	ACTAAAGTAC	GGGGCTCCTT	33000
TGCATGTAAC	AGACGACCTA	AACACTTTGA	CCGTAGCAAC	TGGTCCAGGT	33050
GTGACTATTA	ATAATACTTC	CTTGCAAAC	AAAGTTACTG	GAGCCTTG	33100
TTTTGATTCA	CAAGGCAATA	TGCAACTTAA	TGTAGCAGGA	GGACTAAGGA	33150
TTGATTCTCA	AAACAGACGC	CTTATACTTG	ATGTTAGTTA	TCCGTTTGAT	33200
GCTCAAAACC	AACTAAATCT	AAGACTAGGA	CAGGGCCCTC	TTTTTATAAA	33250
CTCAGCCCAC	AACTTGGATA	TTAACTACAA	CAAAGGCCTT	TACTTGTTTA	33300
CAGCTTCAAA	CAATTCCAAA	AAGCTTGAGG	TTAACCTAAG	CACTGCCAAG	33350
GGGTTGATGT	TTGACGCTAC	AGCCATAGCC	ATTAATGCAG	GAGATGGGCT	33400
TGAATTTGGT	TCACCTAATG	CACCAAACAC	AAATCCCCTC	AAAACAAAAA	33450
TTGGCCATGG	CCTAGAATTT	GATTCAAACA	AGGCTATGGT	TCCTAAACTA	33500
GGAAGTGGCC	TTAGTTTTGA	CAGCACAGGT	GCCATTACAG	TAGGAAACAA	33550
AAATAATGAT	AAGCTAACTT	TGTGGACCAC	ACCAGCTCCA	TCTCCTAACT	33600
GTAGACTAAA	TGCAGAGAAA	GATGCTAAAC	TCACTTTGGT	CTTAACAAAA	33650
TGTGGCAGTC	AAATACTTGC	TACAGTTTCA	GTTTTGGCTG	TTAAAGGCAG	33700
TTTGGCTCCA	ATATCTGGAA	CAGTTCAAAG	TGCTCATCTT	ATTATAAGAT	33750
TTGACGAAAA	TGGAGTGCTA	CTAAACAATT	CCTTCCTGGA	CCCAGAATAT	33800
TGGAACCTTA	GAAATGGAGA	TCTTACTGAA	GGCACAGCCT	ATACAAACGC	33850
TGTTGGATTT	ATGCCTAACC	TATCAGCTTA	TCCAAAATCT	CACGGTAAAA	33900
CTGCCAAAAG	TAACATTGTC	AGTCAAGTTT	ACTTAAACGG	AGACAAAAC	33950
AAACCTGTAA	CACTAACCAT	TACACTAAAC	GGTACACAGG	AAACAGGAGA	34000
CACAACTCCA	AGTGCATACT	CTATGTCATT	TTCATGGGAC	TGGTCTGGCC	34050
ACAACCTACAT	TAATGAAATA	TTTGCCACAT	CCTCTTACAC	TTTTTCATAC	34100

100

ATTGCCCAAG	AATAAAGAAT	CGTTTGTGTT	ATGTTTCAAC	GTGTTTATTT	34150
TTCAATTGCA	GAAAATTTCA	AGTCATTTTT	CATTCAGTAG	TATAGCCCCA	34200
CCACCACATA	GCTTATACAG	ATCACCGTAC	CTTAATCAAA	CTCACAGAAC	34250
CCTAGTATTC	AACCTGCCAC	CTCCCTCCCA	ACACACAGAG	TACACAGTCC	34300
TTTCTCCCCG	GCTGGCCTTA	AAAAGCATCA	TATCATGGGT	AACAGACATA	34350
TTCTTAGGTG	TTATATTCCA	CACGGTTTCC	TGTCGAGCCA	AACGCTCATC	34400
AGTGATATTA	ATAAACTGGC	GGCGATATAA	AATGCAAGGT	GCTGCTCAAA	34450
AAATCAGGCA	AAGCCTCGCG	CAAAAAAGAA	AGCACATCGT	AGTCATGCTC	34500
ATGCAGATAA	AGGCAGGTAA	GCTCCGGAAC	CACCACAGAA	AAAGACACCA	34550
TTTTTCTCTC	AAACATGTCT	GCGGGTTTCT	GCATAAACAC	AAAATAAAAT	34600
AACAAAAAAA	CATTTAAACA	TTAGAAGCCT	GTCTTACAAC	AGGAAAAACA	34650
ACCCTTATAA	GCATAAGACG	GACTACGGCC	ATGCCGGCGT	GACCGTAAAA	34700
AAACTGGTCA	CCGTGATTAA	AAAGCACCAC	CGACAGCTCC	TCGGTCATGT	34750
CCGGAGTCAT	AATGTAAGAC	TCGGTAAACA	CATCAGGTTG	ATTCATCGGT	34800
CAGTGCTAAA	AAGCGACCGA	AATAGCCCGG	GGGAATACAT	ACCCGCAGGC	34850
GTAGAGACAA	CATTACAGCC	CCCATAGGAG	GTATAACAAA	ATTAATAGGA	34900
GAGAAAAACA	CATAAACACC	TGAAAAACCC	TCCTGCCTAG	GCAAAATAGC	34950
ACCCTCCCCG	TCCAGAACAA	CATACAGCGC	TTCACAGCGG	CAGCCTAACA	35000
GTCAGCCTTA	CCAGTAAAAA	AGAAAACCTA	TTAAAAAAAC	ACCACTCGAC	35050
ACGGCACCAG	CTCAATCAGT	CACAGTGTA	AAAAGGGCCA	AGTGCAGAGC	35100
GAGTATATAT	AGGACTAAAA	AATGACGTAA	CGGTTAAAGT	CCACAAAAAA	35150
CACCCAGAAA	ACCGCACGCG	AACCTACGCC	CAGAAACGAA	AGCCAAAAAA	35200
CCCACAACCT	CCTCAAATCG	TCACTTCCGT	TTTCCCACGT	TACGTAACCT	35250
CCCATTTTAA	GAAAACCTACA	ATTCCCAACA	CATACAAGTT	ACTCCGCCCT	35300
AAAACCTACG	TCACCCGCCC	CGTTCCCACG	CCCCGCGCCA	CGTCACAAAC	35350

101

TCCACCCCCT CATTATCATA TTGGCTTCAA TCCAAAATAA GGTATATTAT 35400  
TGATGATG 35408

## (2) INFORMATION FOR SEQ ID NO:4:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 8509 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: not relevant

## (ii) MOLECULE TYPE: cDNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

GCCCAATACG CAAACCGCCT CTCCCCGCGC GTTGGCCGAT TCATTAATGC 50  
AGCTGCGCGC TCGCTCGCTC ACTGAGGCCG CCCGGGCAAA GCCCGGGCGT 100  
CGGGCGACCT TTGGTCGCCC GGCCTCAGTG AGCGAGCGAG CGCGCAGAGA 150  
GGGAGTGGCC AACTCCATCA CTAGGGGTTC CTTGTAGTTA ATGATTAACC 200  
CGCCATGCTA CTTATCTACG TAGCCATTCT CTAGCCCCTG CAGGTCGTTA 250  
CATAACTTAC GGTAAATGGC CCGCCTGGCT GACCGCCCAA CGACCCCCGC 300  
CCATTGACGT CAATAATGAC GTATGTTCCC ATAGTAACGC CAATAGGGAC 350  
TTTCCATTGA CGTCAATGGG TGGAGTATTT ACGGTAAACT GCCCACTTGG 400  
CAGTACATCA AGTGTATCAT ATGCCAAGTA CGCCCCCTAT TGACGTCAAT 450  
GACGGTAAAT GGCCCGCCTG GCATTATGCC CAGTACATGA CCTTATGGGA 500  
CTTTCCTACT TGGCAGTACA TCTACGTATT AGTCATCGCT ATTACCATGG 550  
TGATGCGGTT TTGGCAGTAC ATCAATGGGC GTGGATAGCG GTTTGACTCA 600  
CGGGGATTTC CAAGTCTCCA CCCCATTGAC GTCAATGGGA GTTTGTTTTG 650  
GCACCAAAT CAACGGGACT TTCCAAAATG TCGTAACAAC TCCGCCCCAT 700  
TGACGCAAAT GGGCGGTAGG CGTGTACGGT GGGAGGTCTA TATAAGCAGA 750  
GCTCGTTT TAGAACCGTCA GATCGCCTGG AGACGCCATC CACGCTGTTT 800  
TGACCTCCAT AGAAGACACC GGGACCGATC CAGCCTCCGG ACTCTAGAGG 850  
ATCCGGTACT CGAGGAACTG AAAAACCAGA AAGTTAACTG GTAAGTTTAG 900



102

TCTTTTGTGTC	TTTTATTTCA	GGTCCCGGAT	CCGGTGGTGG	TGCAAATCAA	950	
AGAACTGCTC	CTCAGTGGAT	GTTGCCTTTA	CTTCTAGGCC	TGTACGGAAG	1000	
TGTTACTTCT	GCTCTAAAAG	CTGCGGAATT	GTACCCGCGG	CCGCAATTCC	1050	
CGGGGATCGA	AAGAGCCTGC	TAAAGCAAAA	AAGAAGTCAC	CATGTCGTTT	1100	
ACTTTGACCA	ACAAGAACGT	GATTTTCGTT	GCCGGTCTGG	GAGGCATTGG	1150	
TCTGGACACC	AGCAAGGAGC	TGCTCAAGCG	CGATCCCGTC	GTTTTACAAC	1200	
GTCGTGACTG	GGAAAACCCT	GGCGTTACCC	AACTTAATCG	CCTTGCAGCA	1250	
CATCCCCCTT	TCGCCAGCTG	GCGTAATAGC	GAAGAGGCC	GCACCGATCG	1300	
CCCTTCCCAA	CAGTTGCGCA	GCCTGAATGG	CGAATGGCGC	TTTGCCTGGT	1350	
TTCCGGCACC	AGAAGCGGTG	CCGGAAAGCT	GGCTGGAGTG	CGATCTTCCT	1400	
GAGGCCGATA	CTGTCGTCGT	CCCCTCAAAC	TGGCAGATGC	ACGGTTACGA	1450	
TGCGCCCATC	TACACCAACG	TAACCTATCC	CATTACGGTC	AATCCGCCGT	1500	
TTGTTCCCAC	GGAGAATCCG	ACGGGTGTGT	ACTCGCTCAC	ATTTAATGTT	1550	
GATGAAAGCT	GGCTACAGGA	AGGCCAGACG	CGAATTATTT	TTGATGGCGT	1600	
TAACTCGGCG	TTTCATCTGT	GGTGCAACGG	GCGCTGGGTC	GGTTACGGCC	1650	
AGGACAGTCG	TTTGCCGTCT	GAATTGACC	TGAGCGCATT	TTTACGCGCC	1700	
GGAGAAAACC	GCCTCGCGGT	GATGGTGCTG	CGTTGGAGTG	ACGGCAGTTA	1750	
TCTGGAAGAT	CAGGATATGT	GGCGGATGAG	CGGCATTTTC	CGTGACGTCT	1800	
CGTTGCTGCA	TAAACCGACT	ACACAAATCA	GCGATTTCCA	TGTTGCCACT	1850	
CGCTTTAATG	ATGATTTTCA	CCGCGCTGTA	CTGGAGGCTG	AAGTTCAGAT	1900	
GTGCGGCGAG	TTGCGTGA	ACCTACGGGT	AACAGTTTCT	TTATGGCAGG	1950	
GTGAAACGCA	GGTCGCCAGC	GGCACC	CGCGC	CTTTCGGCGG	TGAAATTATC	2000
GATGAGCGTG	GTGGTTATGC	CGATCGCGTC	ACACTACGTC	TGAACGTCGA	2050	
AAACCCGAAA	CTGTGGAGCG	CCGAAATCCC	GAATCTCTAT	CGTGCGGTGG	2100	
TTGAACTGCA	CACCGCCGAC	GGCAGCTGA	TTGAAGCAGA	AGCCTGCGAT	2150	
GTCGGTTTCC	GCGAGGTGCG	GATTGAAAAT	GGTCTGCTGC	TGCTGAACGG	2200	

103

CAAGCCGTTG	CTGATTTCGAG	GCGTTAACCG	TCACGAGCAT	CATCCTCTGC	2250
ATGGTCAGGT	CATGGATGAG	CAGACGATGG	TGCAGGATAT	CCTGCTGATG	2300
AAGCAGAACA	ACTTTAACGC	CGTGCGCTGT	TCGCATTATC	CGAACCATCC	2350
GCTGTGGTAC	ACGCTGTGCG	ACCGCTACGG	CCTGTATGTG	GTGGATGAAG	2400
CCAATATTGA	AACCCACGGC	ATGGTGCCAA	TGAATCGTCT	GACCGATGAT	2450
CCGCGCTGGC	TACCGGCGAT	GAGCGAACGC	GTAACGCGAA	TGGTGACGCG	2500
CGATCGTAAT	CACCCGAGTG	TGATCATCTG	GTCGCTGGGG	AATGAATCAG	2550
GCCACGGCGC	TAATCACGAC	GCGCTGTATC	GCTGGATCAA	ATCTGTGCGAT	2600
CCTTCCCGCC	CGGTGCAGTA	TGAAGGCGGC	GGAGCCGACA	CCACGGCCAC	2650
CGATATTATT	TGCCCCGATGT	ACGCGCGCGT	GGATGAAGAC	CAGCCCTTCC	2700
CGGCTGTGCC	GAAATGGTCC	ATCAAAAAAT	GGCTTTCGCT	ACCTGGAGAG	2750
ACGCGCCCGC	TGATCCTTTG	CGAATACGCC	CACGCGATGG	GTAACAGTCT	2800
TGGCGGTTTC	GCTAAATACT	GGCAGGCGTT	TCGTCAGTAT	CCCCGTTTAC	2850
AGGGCGGCTT	CGTCTGGGAC	TGGGTGGATC	AGTCGCTGAT	TAAATATGAT	2900
GAAAACGGCA	ACCCGTGGTC	GGCTTACGGC	GGTGATTTTG	GCGATACGCC	2950
GAACGATCGC	CAGTTCTGTA	TGAACGGTCT	GGTCTTTGCC	GACCGCACGC	3000
CGCATCCAGC	GCTGACGGAA	GCAAAACACC	AGCAGCAGTT	TTTCCAGTTC	3050
CGTTTATCCG	GGCAAACCAT	CGAAGTGACC	AGCGAATACC	TGTTCCGTCA	3100
TAGCGATAAC	GAGCTCCTGC	ACTGGATGGT	GGCGCTGGAT	GGTAAGCCGC	3150
TGGCAAGCGG	TGAAGTGCCT	CTGGATGTGG	CTCCACAAGG	TAAACAGTTG	3200
ATTGAACTGC	CTGAACTACC	GCAGCCGGAG	AGCGCCGGGC	AACTCTGGCT	3250
CACAGTACGC	GTAGTGCAAC	CGAACGCGAC	CGCATGGTCA	GAAGCCGGGC	3300
ACATCAGCGC	CTGGCAGCAG	TGGCGTCTGG	CGGAAAACCT	CAGTGTGACG	3350
CTCCCCGCCG	CGTCCCACGC	CATCCCGCAT	CTGACCACCA	GCGAAATGGA	3400
TTTTTGCATC	GAGCTGGGTA	ATAAGCGTTG	GCAATTTAAC	CGCCAGTCAG	3450
GCTTTCCTTC	ACAGATGTGG	ATTGGCGATA	AAAAACAAC	GCTGACGCCG	3500

CTGCGCGATC	AGTTCACCCG	TGCACCGCTG	GATAACGACA	TTGGCGTAAG	3550
TGAAGCGACC	CGCATTGACC	CTAACGCCTG	GGTCGAACGC	TGGAAGGCGG	3600
CGGGCCATTA	CCAGGCCGAA	GCAGCGTTGT	TGCAGTGCAC	GGCAGATACA	3650
CTTGCTGATG	CGGTGCTGAT	TACGACCGCT	CACGCGTGGC	AGCATCAGGG	3700
GAAAACCTTA	TTTATCAGCC	GGAAAACCTA	CCGGATTGAT	GGTAGTGGTC	3750
AAATGGCGAT	TACCGTTGAT	GTTGAAGTGG	CGAGCGATAC	ACCGCATCCG	3800
GCGCGGATTG	GCCTGAACTG	CCAGCTGGCG	CAGGTAGCAG	AGCGGGTAAA	3850
CTGGCTCGGA	TTAGGGCCGC	AAGAAAACCTA	TCCCGACCGC	CTTACTGCCC	3900
CCTGTTTTGA	CCGCTGGGAT	CTGCCATTGT	CAGACATGTA	TACCCCGTAC	3950
GTCTTCCCGA	GCGAAAACGG	TCTGCGCTGC	GGGACGCGCG	AATTGAATTA	4000
TGGCCACAC	CAGTGGCGCG	GCGACTTCCA	GTTCAACATC	AGCCGCTACA	4050
GTCAACAGCA	ACTGATGGAA	ACCAGCCATC	GCCATCTGCT	GCACGCGGAA	4100
GAAGGCACAT	GGCTGAATAT	CGACGGTTTC	CATATGGGGA	TTGGTGGCGA	4150
CGACTCCTGG	AGCCCGTCAG	TATCGGCGGA	ATTACAGCTG	AGCGCCGGTC	4200
GCTACCATTA	CCAGTTGGTC	TGGTGTCAAA	AATAATAATA	ACCGGGCAGG	4250
CCATGTCTGC	CCGTATTTTCG	CGTAAGGAAA	TCCATTATGT	ACTATTTAAA	4300
AAACACAAAC	TTTTGGATGT	TCGGTTTATT	CTTTTTCTTT	TACTTTTTTA	4350
TCATGGGAGC	CTACTTCCCG	TTTTTCCCGA	TTTGGCTACA	TGACATCAAC	4400
CATATCAGCA	AAAGTGATAC	GGGTATTATT	TTTGCCGCTA	TTTCTCTGTT	4450
CTCGCTATTA	TTCCAACCGC	TGTTTGGTCT	GCTTTCTGAC	AAACTCGGCC	4500
TCGACTCTAG	GCGGCCGCGG	GGATCCAGAC	ATGATAAGAT	ACATTGATGA	4550
GTTTGGACAA	ACCACAACCTA	GAATGCAGTG	AAAAAAATGC	TTTATTTGTG	4600
AAATTTGTGA	TGCTATTGCT	TTATTTGTAA	CCATTATAAG	CTGCAATAAA	4650
CAAGTTAACA	ACAACAATTG	CATTCATTTT	ATGTTTCAGG	TTCAGGGGGA	4700
GGTGTGGGAG	GTTTTTTCGG	ATCCTCTAGA	GTCGACCTGC	AGGGGCTAGA	4750
ATGGCTACGT	AGATAAGTAG	CATGGCGGGT	TAATCATTA	CTACAAGGAA	4800

105

CCCCTAGTGA	TGGAGTTGGC	CACTCCCTCT	CTGCGCGCTC	GCTCGCTCAC	4850
TGAGGCCGGG	CGACCAAAGG	TCGCCCACG	CCCGGGCTTT	GCCCGGGCGG	4900
CCTCAGTGAG	CGAGCGAGCG	CGCAGCTGGC	GTAATAGCGA	AGAGGCCCCG	4950
ACCGATCGCC	CTTCCCAACA	GTTGCGCAGC	CTGAATGGCG	AATGGAATTC	5000
CAGACGATTG	AGCGTCAAAA	TGTAGGTATT	TCCATGAGCG	TTTTTCCTGT	5050
TGCAATGGCT	GGCGGTAATA	TTGTTCTGGA	TATTACCAGC	AAGGCCGATA	5100
GTTTGAGTTC	TTCTACTCAG	GCAAGTGATG	TTATTACTAA	TCAAAGAAGT	5150
ATTGCGACAA	CGGTTAATTT	GCGTGATGGA	CAGACTCTTT	TACTCGGTGG	5200
CCTCACTGAT	TATAAAAACA	CTTCTCAGGA	TTCTGGCGTA	CCGTTCCCTGT	5250
CTAAAATCCC	TTTAATCGGC	CTCCTGTTTA	GCTCCCGCTC	TGATTCTAAC	5300
GAGGAAAGCA	CGTTATACGT	GCTCGTCAAA	GCAACCATAG	TACGCGCCCT	5350
GTAGCGGCGC	ATTAAGCGCG	GCGGGTGTGG	TGGTTACGCG	CAGCGTGACC	5400
GCTACACTTG	CCAGCGCCCT	AGCGCCCGCT	CCTTTCGCTT	TCTTCCCTTC	5450
CTTTCCTCGC	ACGTTGCGCG	GCTTTCCTCG	TCAAGCTCTA	AATCGGGGGC	5500
TCCCTTTAGG	GTTCCGATTT	AGTGCTTTAC	GGCACCTCGA	CCCCAAAAA	5550
CTTGATTAGG	GTGATGGTTC	ACGTAGTGGG	CCATCGCCCT	GATAGACGGT	5600
TTTTCGCCCT	TTGACGTTGG	AGTCCACGTT	CTTTAATAGT	GGACTCTTGT	5650
TCCAAACTGG	AACAACACTC	AACCCTATCT	CGGTCTATT	TTTTGATTTA	5700
TAAGGGATTT	TGCCGATTT	GGCCTATTGG	TTAAAAAATG	AGCTGATTTA	5750
ACAAAAATTT	AACGCGAATT	TTAACAAAAT	ATTAACGTTT	ACAATTTAAA	5800
TATTTGCTTA	TACAATCTTC	CTGTTTTTGG	GGCTTTTCTG	ATTATCAACC	5850
GGGGTACATA	TGATTGACAT	GCTAGTTTTA	CGATTACCGT	TCATCGATTC	5900
TCTTGTTTGC	TCCAGACTCT	CAGGCAATGA	CCTGATAGCC	TTTGTAGAGA	5950
CCTCTCAAAA	ATAGCTACCC	TCTCCGGCAT	GAATTTATCA	GCTAGAACGG	6000
TTGAATATCA	TATTGATGGT	GATTTGACTG	TCTCCGGCCT	TTCTCACCCG	6050
TTTGAATCTT	TACCTACACA	TTACTCAGGC	ATTGCATTTA	AAATATATGA	6100

GGGTTCTAAA	AATTTTTTATC	CTTGCGTTGA	AATAAAGGCT	TCTCCCGCAA	6150
AAGTATTACA	GGGTCATAAT	GTTTTTGGA	CAACCGATTT	AGCTTTATGC	6200
TCTGAGGCTT	TATTGCTTAA	TTTTGCTAAT	TCTTTGCCTT	GCCTGTATGA	6250
TTTATTGGAT	GTTGGAATTC	CTGATGCGGT	ATTTTCTCCT	TACGCATCTG	6300
TGCGGTATTT	CACACCGCAT	ATGGTGCACT	CTCAGTACAA	TCTGCTCTGA	6350
TGCCGCATAG	TTAAGCCAGC	CCCGACACCC	GCCAACACCC	GCTGACGCGC	6400
CCTGACGGGC	TTGTCTGCTC	CCGGCATCCG	CTTACAGACA	AGCTGTGACC	6450
GTCTCCGGGA	GCTGCATGTG	TCAGAGGTTT	TCACCGTCAT	CACCGAAACG	6500
CGCGAGACGA	AAGGGCCTCG	TGATACGCCT	ATTTTTATAG	GTTAATGTCA	6550
TGATAATAAT	GGTTTCTTAG	ACGTCAGGTG	GCACTTTTCG	GGGAAATGTG	6600
CGCGGAACCC	CTATTTGTTT	ATTTTTCTAA	ATACATTCAA	ATATGTATCC	6650
GCTCATGAGA	CAATAACCCT	GATAAATGCT	TCAATAATAT	TGAAAAAGGA	6700
AGAGTATGAG	TATTCAACAT	TTCCGTGTG	CCCTTATTCC	CTTTTTTGCG	6750
GCATTTTGCC	TTCCTGTTTT	TGCTCACCCA	GAAACGCTGG	TGAAAGTAAA	6800
AGATGCTGAA	GATCAGTTGG	GTGCACGAGT	GGGTACATC	GAAGTGGATC	6850
TCAACAGCGG	TAAGATCCTT	GAGAGTTTTT	GCCCCGAAGA	ACGTTTTCCA	6900
ATGATGAGCA	CTTTTAAAGT	TCTGCTATGT	GGCGCGGTAT	TATCCCGTAT	6950
TGACGCCGGG	CAAGAGCAAC	TCGGTCGCCG	CATACACTAT	TCTCAGAATG	7000
ACTTGGTTGA	GTAATCACCA	GTCACAGAAA	AGCATCTTAC	GGATGGCATG	7050
ACAGTAAGAG	AATTATGCAG	TGCTGCCATA	ACCATGAGTG	ATAACACTGC	7100
GGCCAACTTA	CTTCTGACAA	CGATCGGAGG	ACCGAAGGAG	CTAACCGCTT	7150
TTTTGCACAA	CATGGGGGAT	CATGTAATC	GCCTTGATCG	TTGGGAACCG	7200
GAGCTGAATG	AAGCCATACC	AAACGACGAG	CGTGACACCA	CGATGCCTGT	7250
AGCAATGGCA	ACAACGTTGC	GCAAATATT	AACTGGCGAA	CTACTTACTC	7300
TAGCTTCCCG	GCAACAATTA	ATAGACTGGA	TGGAGGCGGA	TAAAGTTGCA	7350
GGACCACTTC	TGCGCTCGGC	CCTTCCGGCT	GGCTGGTTTA	TTGCTGATAA	7400

107

ATCTGGAGCC	GGTGAGCGTG	GGTCTCGCGG	TATCATTGCA	GCACTGGGGC	7450
CAGATGGTAA	GCCCTCCCGT	ATCGTAGTTA	TCTACACGAC	GGGGAGTCAG	7500
GCAACTATGG	ATGAACGAAA	TAGACAGATC	GCTGAGATAG	GTGCCTCACT	7550
GATTAAGCAT	TGGTAACTGT	CAGACCAAGT	TTACTCATAT	ATACTTTAGA	7600
TTGATTTAAA	ACTTCATTTT	TAATTTAAAA	GGATCTAGGT	GAAGATCCTT	7650
TTTGATAATC	TCATGACCAA	AATCCCTTAA	CGTGAGTTTT	CGTTCCACTG	7700
AGCGTCAGAC	CCCGTAGAAA	AGATCAAAGG	ATCTTCTTGA	GATCCTTTTT	7750
TTCTGCGCGT	AATCTGCTGC	TTGCAAACAA	AAAAACCACC	GCTACCAGCG	7800
GTGGTTTGTT	TGCCGGATCA	AGAGCTACCA	ACTCTTTTTC	CGAAGGTAAC	7850
TGGCTTCAGC	AGAGCGCAGA	TACCAAATAC	TGTCCTTCTA	GTGTAGCCGT	7900
AGTTAGGCCA	CCACTTCAAG	AACTCTGTAG	CACCGCCTAC	ATACCTCGCT	7950
CTGCTAATCC	TGTTACCAGT	GGCTGCTGCC	AGTGGCGATA	AGTCGTGTCT	8000
TACCGGGTTG	GACTCAAGAC	GATAGTTACC	GGATAAGGCG	CAGCGGTCGG	8050
GCTGAACGGG	GGGTTCGTGC	ACACAGCCCA	GCTTGGAGCG	AACGACCTAC	8100
ACCGAACTGA	GATACCTACA	GCGTGAGCTA	TGAGAAAGCG	CCACGCTTCC	8150
CGAAGGGAGA	AAGGCGGACA	GGTATCCGGT	AAGCGGCAGG	GTCGGAACAG	8200
GAGAGCGCAC	GAGGGAGCTT	CCAGGGGGAA	ACGCCTGGTA	TCTTTATAGT	8250
CCTGTCGGGT	TTCGCCACCT	CTGACTTGAG	CGTCGATTTT	TGTGATGCTC	8300
GTCAGGGGGG	CGGAGCCTAT	GGAAAAACGC	CAGCAACGCG	GCCTTTTAC	8350
GGTTCCTGGC	CTTTTGCTGG	CCTTTTGCTC	ACATGTTCTT	TCCTGCGTTA	8400
TCCCCTGATT	CTGTGGATAA	CCGTATTACC	GCCTTTGAGT	GAGCTGATAC	8450
CGCTCGCCGC	AGCCGAACGA	CCGAGCGCAG	CGAGTCAGTG	AGCGAGGAAG	8500
CGGAAGAGC					8509

108

## (2) INFORMATION FOR SEQ ID NO:5:

## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 8299 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: not relevant

## (ii) MOLECULE TYPE: cDNA

## (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

GCCCAATACG CAAACCGCCT CTCCCCGCGC GTTGGCCGAT TCATTAATGC	50
AGCTGCGCGC TCGCTCGCTC ACTGAGGCCG CCCGGGCAAA GCCCGGGCGT	100
CGGGCGACCT TTGGTCGCCC GGCCTCAGTG AGCGAGCGAG CGCGCAGAGA	150
GGGAGTGGCC AACTCCATCA CTAGGGGTTC CTTGTAGTTA ATGATTAACC	200
CGCCATGCTA CTTATCTACA TCATCGATGA ATTCGAGCTT GCATGCCTGC	250
AGGTCGTTAC ATAACCTACG GTAAATGGCC CGCCTGGCTG ACCGCCCCAAC	300
GACCCCCGCC CATTGACGTC AATAATGACG TATGTTCCCA TAGTAACGCC	350
AATAGGGACT TTCCATTGAC GTCAATGGGT GGAGTATTTA CGGTAAACTG	400
CCCACTTGGC AGTACATCAA GTGTATCATA TGCCAAGTAC GCCCCCTATT	450
GACGTCAATG ACGGTAAATG GCCCGCCTGG CATTATGCCC AGTACATGAC	500
CTTATGGGAC TTTCTACTT GGCAGTACAT CTACGTATTA GTCATCGCTA	550
TTACCATGGT GATGCGGTTT TGGCAGTACA TCAATGGGCG TGGATAGCGG	600
TTTGACTCAC GGGGATTTC AAGTCTCCAC CCCATTGACG TCAATGGGAG	650
TTTGTTTTTG CACCAAAATC AACGGGACTT TCCAAAATGT CGTAACAACT	700
CCGCCCCATT GACGCAAATG GGCGGTAGGC GTGTACGGTG GGAGGTCTAT	750
ATAAGCAGAG CTCGTTTAGT GAACCGTCAG ATCGCCTGGA GACGCCATCC	800
ACGCTGTTTT GACCTCCATA GAAGACACCG GGACCGATCC AGCCTCCGGA	850
CTCTAGAGGA TCCGGTACTC GACCCGAGCT CGGATCCACT AGTAACGGCC	900
GCCAGTGTGC TGGAATTCTG CACTCCAGGC TGCCCGGGTT TGCATGCTGC	950
TGCTGCTGCT GCTGCTGGGC CTGAGGCTAC AGCTCTCCCT GGGCATCATC	1000

109

CTAGTTGAGG	AGGAGAACCC	GGACTTCTGG	AACCGCGAGG	CAGCCGAGGC	1050
CCTGGGTGCC	GCCAAGAAGC	TGCAGCCTGC	ACAGACAGCC	GCCAAGAACC	1100
TCATCATCTT	CCTGGGCGAT	GGGATGGGGG	TGTCTACGGT	GACAGCTGCC	1150
AGGATCCTAA	AAGGGCAGAA	GAAGGACAAA	CTGGGGCCTG	AGATACCCCT	1200
GGCCATGGAC	CGCTTCCCAT	ATGTGGCTCT	GTCCAAGACA	TACAATGTAG	1250
ACAAACATGT	GCCAGACAGT	GGAGCCACAG	CCACGGCCTA	CCTGTGCGGG	1300
GTCAAGGGCA	ACTTCCAGAC	CATTGGCTTG	AGTGCAGCCG	CCCGCTTTAA	1350
CCAGTGCAAC	ACGACACGCG	GCAACGAGGT	CATCTCCGTG	ATGAATCGGG	1400
CCAAGAAAGC	AGGGAAGTCA	GTGGGAGTGG	TAACCACCAC	ACGAGTGCAG	1450
CACGCCTCGC	CAGCCGGCAC	CTACGCCCAC	ACGGTGAACC	GCAACTGGTA	1500
CTCGGACGCC	GACGTGCCTG	CCTCGGCCCC	CCAGGAGGGG	TGCCAGGACA	1550
TCGCTACGCA	GCTCATCTCC	AACATGGACA	TTGATGTGAT	CCTAGGTGGA	1600
GGCCGAAAGT	ACATGTTTCG	CATGGGAACC	CCAGACCCTG	AGTACCCAGA	1650
TGACTACAGC	CAAGGTGGGA	CCAGGCTGGA	CGGGAAGAAT	CTGGTGCAGG	1700
AATGGCTCGG	CGAACGCCAG	GGTGCCCGGT	ACGTGTGGAA	CCGCACTGAG	1750
CTCATGCAGG	CTTCCCTGGA	CCCGTCTGTG	ACCCATCTCA	TGGGTCTCTT	1800
TGAGCCTGGA	GACATGAAAT	ACGAGATCCA	CCGAGACTCC	ACACTGGACC	1850
CCTCCCTGAT	GGAGATGACA	GAGGCTGCCC	TGCGCCTGCT	GAGCAGACAC	1900
CCCCGCGGCT	TCTTCCTCTT	CGTGGAGGGT	GGTCGCATCG	ACCATGGTCA	1950
TCATGAAAGC	AGGGCTTACC	GGGCACTGAC	TGAGACGATC	ATGTTTCGACG	2000
ACGCCATTGA	GAGGGCGGGC	CAGCTCACCA	GCGAGGAGGA	CACGCTGAGC	2050
CTCGTCACTG	CCGACCACTC	CCACGTCTTC	TCCTTCGGAG	GCTACCCCT	2100
GCGAGGGAGC	TCCTTCATCG	GGCTGGCCGC	TGGCAAGGCC	CGGGACAGGA	2150
AGGCCTACAC	GGTCCTCCTA	TACGGAAACG	GTCCAGGCTA	TGTGCTCAAG	2200
GACGGCGCCC	GGCCGGATGT	TACCGAGAGC	GAGAGCGGGA	GCCCCGAGTA	2250
TCGGCAGCAG	TCAGCAGTGC	CCCTGGACGA	AGAGACCCAC	GCAGGCGAGG	2300



110

ACGTGGCGGT	GTTCGCGCGC	GGCCCGCAGG	CGCACCTGGT	TCACGGCGTG	2350
CAGGAGCAGA	CCTTCATAGC	GCACGTCATG	GCCTTCGCCG	CCTGCCTGGA	2400
GCCCTACACC	GCCTGCGACC	TGGCGCCCCC	CGCCGGCACC	ACCGACGCCG	2450
CGCACCCGGG	GCGGTCCGTG	GTCCCCGCGT	TGCTTCCTCT	GCTGGCCGGG	2500
ACCCTGCTGC	TGCTGGAGAC	GGCCACTGCT	CCCTGAGTGT	CCCGTCCCTG	2550
GGGCTCCTGC	TTCCCCATCC	CGGAGTTCTC	CTGCTCCCCA	CCTCCTGTCTG	2600
TCCTGCCTGG	CCTCCAGCCC	GAGTCGTCAT	CCCCGGAGTC	CCTATACAGA	2650
GGTCCTGCCA	TGGAACCTTC	CCCTCCCCGT	GCGCTCTGGG	GACTGAGCCC	2700
ATGACACCAA	ACCTGCCCCCT	TGGCTGCTCT	CGGACTCCCT	ACCCCAACCC	2750
CAGGGACTGC	AGGTTGTGCC	CTGTGGCTGC	CTGCACCCCA	GGAAAGGAGG	2800
GGGCTCAGGC	CATCCAGCCA	CCACCTACAG	CCCAGTGGGG	TCGAGACAGA	2850
TGGTCAGTCT	GGAGGATGAC	GTGGCGTGAA	GCTGGCCGCG	GGGATCCAGA	2900
CATGATAAGA	TACATTGATG	AGTTTGGACA	AACCACAACT	AGAATGCAGT	2950
GAAAAAATG	CTTTATTTGT	GAAATTTGTG	ATGCTATTGC	TTTATTTGTA	3000
ACCATTATAA	GCTGCAATAA	ACAAGTTAAC	AACAACAATT	GCATTCATTT	3050
TATGTTTCAG	GTTCAGGGGG	AGGTGTGGGA	GGTTTTTTTCG	GATCCTCTAG	3100
AGTCGACTCT	AGANNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	3150
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	3200
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	3250
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	3300
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	3350
NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNNNNNNNN	NNNGGATCCC	3400
CATGACTACG	TCCGGCGTTC	CATTTGGCAT	GACACTACGA	CCAACACGAT	3450
CTCGGTTGTC	TCGGCGCACT	CCGTACAGTA	GGGATCGTCT	ACCTCCTTTT	3500
GAGACAGAAA	CCCGCGCTAC	CATACTGGAG	GATCATCCGC	TGCTGCCCCG	3550
ATGTAACACT	TTGACAATGC	ACAACGTGAG	TTACGTGCGA	GGTCTTCCCT	3600

111

GCAGTGTGGG	ATTTACGCTG	ATTCAGGAAT	GGGTTGTTCC	CTGGGATATG	3650
GTTCTAACGC	GGGAGGAGCT	TGTAATCCTG	AGGAAGTGTA	TGCACGTGTG	3700
CCTGTGTTGT	GCCAACATTG	ATATCATGAC	GAGCATGATG	ATCCATGGTT	3750
ACGAGTCCTG	GGCTCTCCAC	TGTCATTGTT	CCAGTCCCGG	TTCCCTGCAG	3800
TGTATAGCCG	GCGGGCAGGT	TTTGGCCAGC	TGGTTTAGGA	TGGTGGTGGA	3850
TGGCGCCATG	TTTAATCAGA	GGTTTATATG	GTACCGGGAG	GTGGTGAATT	3900
ACAACATGCC	AAAAGAGGTA	ATGTTTATGT	CCAGCGTGTT	TATGAGGGGT	3950
CGCCACTTAA	TCTACCTGCG	CTTGTGGTAT	GATGGCCACG	TGGGTTCTGT	4000
GGTCCCCGCC	ATGAGCTTTG	GATACAGCGC	CTTGCACTGT	GGGATTTTGA	4050
ACAATATTGT	GGTGCTGTGC	TGCAGTTACT	GTGCTGATTT	AAGTGAGATC	4100
AGGGTGCGCT	GCTGTGCCCC	GAGGACAAGG	CGCCTTATGC	TGCGGGCGGT	4150
GCGAATCATC	GCTGAGGAGA	CCACTGCCAT	GTTGTATTCC	TGCAGGACGG	4200
AGCGGCGGCG	GCAGCAGTTT	ATTCGCGCGC	TGCTGCAGCA	CCACCGCCCT	4250
ATCCTGATGC	ACGATTATGA	CTCTACCCCC	ATGTAGGGAT	CCCCATCACT	4300
AGTGCGGCGG	CGGGGATCCA	GACATGATAA	GATACATTGA	TGAGTTTGGA	4350
CAAACCACAA	CTAGAATGCA	GTGAAAAAAA	TGCTTTATTT	GTGAAATTTG	4400
TGATGCTATT	GCTTTATTTG	TAACCATTAT	AAGCTGCAAT	AAACAAGTTA	4450
ACAACAACAA	TTGCATTTCAT	TTTATGTTTC	AGGTTCAGGG	GGAGGTGTGG	4500
GAGGTTTTTT	CGGATCCTCT	AGAGTCGACC	TGCAGGCATG	CAAGCTGTAG	4550
ATAAGTAGCA	TGGCGGGTTA	ATCATTAACT	ACAAGGAACC	CCTAGTGATG	4600
GAGTTGGCCA	CTCCCTCTCT	GCGCGCTCGC	TCGCTCACTG	AGGCCGGGCG	4650
ACCAAAGGTC	GCCCGACGCC	CGGGCTTTGC	CCGGGCGGCC	TCAGTGAGCG	4700
AGCGAGCGCG	CAGCTGGCGT	AATAGCGAAG	AGGCCCGCAC	CGATCGCCCT	4750
TCCCAACAGT	TGCGCAGCCT	GAATGGCGAA	TGGAANTTCC	AGACGATTGA	4800
GCGTCAAAAT	GTAGGTATTT	CCATGAGCGT	TTTTCTGT	GCAATGGCTG	4850
GCGGTAATAT	TGTTCTGGAT	ATTACCAGCA	AGGCCGATAG	TTTGAGTTCT	4900

112

TCTACTCAGG	CAAGTGATGT	TATTACTAAT	CAAAGAAGTA	TTGCGACAAC	4950
GGTTAATTTG	CGTGATGGAC	AGACTCTTTT	ACTCGGTGGC	CTCACTGATT	5000
ATAAAAACAC	TTCTCAGGAT	TCTGGCGTAC	CGTTCCTGTC	TAAAATCCCT	5050
TTAATCGGCC	TCCTGTTTAG	CTCCCGCTCT	GATTCTAACG	AGGAAAGCAC	5100
GTTATACGTG	CTCGTCAAAG	CAACCATAGT	ACGCGCCCTG	TAGCGGCGCA	5150
TTAAGCGCGG	CGGGTGTGGT	GGTTACGCGC	AGCGTGACCG	CTACACTTGC	5200
CAGCGCCCTA	GCGCCCGCTC	CTTTCGCTTT	CTTCCCTTCC	TTTCTCGCCA	5250
CGTTCGCCGG	CTTTCCCCGT	CAAGCTCTAA	ATCGGGGGCT	CCCTTTAGGG	5300
TTCCGATTTA	GTGCTTTACG	GCACCTCGAC	CCCAAAAAAC	TTGATTAGGG	5350
TGATGGTTCA	CGTAGTGGGC	CATCGCCCTG	ATAGACGGTT	TTTCGCCCTT	5400
TGACGTTGGA	GTCCACGTTT	TTTAATAGTG	GA CTCTTGTT	CCAAACTGGA	5450
ACAACACTCA	ACCCTATCTC	GGTCTATTCT	TTTGATTAT	AAGGGATTTT	5500
GCCGATTTCTG	GCCTATTGGT	TAAAAAATGA	GCTGATTTAA	CAAAAATTTA	5550
ACGCGAATTT	TAACAAAATA	TTAACGTTTA	CAATTTAAAT	ATTTGCTTAT	5600
ACAATCTTCC	TGTTTTTGGG	GCTTTTCTGA	TTATCAACCG	GGGTACATAT	5650
GATTGACATG	CTAGTTTAC	GATTACCGTT	CATCGATTCT	CTTGTTTGCT	5700
CCAGACTCTC	AGGCAATGAC	CTGATAGCCT	TTGTAGAGAC	CTCTCAAAAA	5750
TAGCTACCCT	CTCCGGCATG	AATTTATCAG	CTAGAACGGT	TGAATATCAT	5800
ATTGATGGTG	ATTTGACTGT	CTCCGGCCTT	TCTCACCCGT	TTGAATCTTT	5850
ACCTACACAT	TACTCAGGCA	TTGCATTTAA	AATATATGAG	GGTTCTAAAA	5900
ATTTTTATCC	TTGCGTTGAA	ATAAAGGCTT	CTCCCGCAAA	AGTATTACAG	5950
GGTCATAATG	TTTTTGGTAC	AACCGATTTA	GCTTTATGCT	CTGAGGCTTT	6000
ATTGCTTAAT	TTTGCTAATT	CTTTGCCTTG	CCTGTATGAT	TTATTGGATG	6050
TTGGAANTTC	CTGATGCGGT	ATTTTCTCCT	TACGCATCTG	TGCGGTATTT	6100
CACACCGCAT	ATGGTGCACT	CTCAGTACAA	TCTGCTCTGA	TGCCGCATAG	6150
TTAAGCCAGC	CCCACACCC	GCCAACACCC	GCTGACGCGC	CCTGACGGGC	6200

113

TTGTCTGCTC	CCGGCATCCG	CTTACAGACA	AGCTGTGACC	GTCTCCGGGA	6250
GCTGCATGTG	TCAGAGGTTT	TCACCGTCAT	CACCGAAACG	CGCGAGACGA	6300
AAGGGCCTCG	TGATACGCCT	ATTTTTATAG	GTTAATGTCA	TGATAATAAT	6350
GGTTTCTTAG	ACGTCAGGTG	GCACTTTTCG	GGGAAATGTG	CGCGGAACCC	6400
CTATTTGTTT	ATTTTTCTAA	ATACATTCAA	ATATGTATCC	GCTCATGAGA	6450
CAATAACCCCT	GATAAATGCT	TCAATAATAT	TGAAAAAGGA	AGAGTATGAG	6500
TATTCACAT	TTCCGTGTCG	CCCTTATTCC	CTTTTTTGCG	GCATTTTGCC	6550
TTCCTGTTTT	TGCTCACCCA	GAAACGCTGG	TGAAAGTAAA	AGATGCTGAA	6600
GATCAGTTGG	GTGCACGAGT	GGGTACATC	GAACGGATC	TCAACAGCGG	6650
TAAGATCCTT	GAGAGTTTTT	GCCCCGAAGA	ACGTTTTCCA	ATGATGAGCA	6700
CTTTTAAAGT	TCTGCTATGT	GGCGCGGTAT	TATCCCGTAT	TGACGCCGGG	6750
CAAGAGCAAC	TCGGTCGCCG	CATACACTAT	TCTCAGAATG	ACTTGTTGA	6800
GTACTCACCA	GTCACAGAAA	AGCATCTTAC	GGATGGCATG	ACAGTAAGAG	6850
AATTATGCAG	TGCTGCCATA	ACCATGAGTG	ATAAACTGC	GGCCAACTTA	6900
CTTCTGACAA	CGATCGGAGG	ACCGAAGGAG	CTAACCGCTT	TTTTGCACAA	6950
CATGGGGGAT	CATGTAAGTC	GCCTTGATCG	TTGGGAACCG	GAGCTGAATG	7000
AAGCCATACC	AAACGACGAG	CGTGACACCA	CGATGCCTGT	AGCAATGGCA	7050
ACAACGTTGC	GCAAACCTATT	AACTGGCGAA	CTACTTACTC	TAGCTTCCCC	7100
GCAACAATTA	ATAGACTGGA	TGGAGGCGGA	TAAAGTTGCA	GGACCACTTC	7150
TGCGCTCGGC	CCTTCCGGCT	GGCTGGTTTA	TTGCTGATAA	ATCTGGAGCC	7200
GGTGAGCGTG	GGTCTCGCGG	TATCATTGCA	GCACTGGGGC	CAGATGGTAA	7250
GCCCTCCCGT	ATCGTAGTTA	TCTACACGAC	GGGGAGTCAG	GCAACTATGG	7300
ATGAACGAAA	TAGACAGATC	GCTGAGATAG	GTGCCTCACT	GATTAAGCAT	7350
TGGTAACTGT	CAGACCAAGT	TTACTCATAT	ATACTTTAGA	TTGATTTAAA	7400
ACTTCATTTT	TAATTTAAAA	GGATCTAGGT	GAAGATCCTT	TTTGATAATC	7450
TCATGACCAA	AATCCCTTAA	CGTGAGTTTT	CGTTCCACTG	AGCGTCAGAC	7500

CCCGTAGAAA	AGATCAAAGG	ATCTTCTTGA	GATCCTTTTT	TTCTGCGCGT	7550
AATCTGCTGC	TTGCAAACAA	AAAAACCACC	GCTACCAGCG	GTGGTTTGTT	7600
TGCCGGATCA	AGAGCTACCA	ACTCTTTTTT	CGAAGGTAAC	TGGCTTCAGC	7650
AGAGCGCAGA	TACCAAATAC	TGTCCTTCTA	GTGTAGCCGT	AGTTAGGCCA	7700
CCACTTCAAG	AACTCTGTAG	CACCGCCTAC	ATACCTCGCT	CTGCTAATCC	7750
TGTTACCAGT	GGCTGCTGCC	AGTGGCGATA	AGTCGTGTCT	TACCGGGTTG	7800
GACTCAAGAC	GATAGTTACC	GGATAAGGCG	CAGCGGTCGG	GCTGAACGGG	7850
GGGTTCGTGC	ACACAGCCCA	GCTTGGAGCG	AACGACCTAC	ACCGAACTGA	7900
GATACCTACA	GCGTGAGCTA	TGAGAAAGCG	CCACGCTTCC	CGAAGGGAGA	7950
AAGGCGGACA	GGTATCCGGT	AAGCGGCAGG	GTCGGAACAG	GAGAGCGCAC	8000
GAGGGAGCTT	CCAGGGGGAA	ACGCCTGGTA	TCTTTATAGT	CCTGTCGGGT	8050
TTCGCCACCT	CTGACTTGAG	CGTCGATTTT	TGTGATGCTC	GTCAGGGGGG	8100
CGGAGCCTAT	GGAAAAACGC	CAGCAACGCG	GCCTTTTTTAC	GGTTCCTGGC	8150
CTTTTGCTGG	CCTTTTGCTC	ACATGTTCTT	TCCTGCGTTA	TCCCCTGATT	8200
CTGTGGATAA	CCGTATTACC	GCCTTTGAGT	GAGCTGATAC	CGCTCGCCGC	8250
AGCCGAACGA	CCGAGCGCAG	CGAGTCAGTG	AGCGAGGAAG	CGGAAGAGC	8299

## WHAT IS CLAIMED IS:

1. A method for enhancing the efficiency of transduction of a recombinant AAV into a target cell *ex vivo* comprising the steps of providing a recombinant adeno-associated virus comprising: (a) the DNA of at least a portion of the genome of an adeno-associated virus which portion is capable of transducing a selected gene into a target cell in the absence of cell division; and (b) a selected gene operatively linked to regulatory sequences directing its expression, said gene flanked by the DNA of (a) and capable of expression in the target cell; infecting said target cells with said recombinant adeno-associated virus; contacting said infected cells with an agent which facilitates the conversion of said ss recombinant virus to its double stranded form, wherein said conversion occurs in said target cell, resulting in enhanced transduction of said recombinant virus into said target cell.

2. The method according to claim 1 wherein said agent is a helper virus comprising a selected gene encoding a polypeptide which can enhance the conversion of said single-stranded recombinant virus to double-stranded recombinant virus or functional fragment thereof, said helper virus capable of transducing said selected gene into the target cell in the absence of cell division.

3. The method according to claim 2 wherein said contacting step comprises co-infecting said target cell with said helper virus.

4. The method according to claim 3 wherein said helper virus is an adenovirus.

5. The method according to claim 2 wherein said selected gene is the adenovirus E4 gene or a functional fragment thereof.

6. The method according to claim 2 wherein said adenovirus contains two selected genes or functional fragments thereof, said genes being the adenovirus E4 gene and E1 gene.

7. The method according to claim 5 or 6 wherein said functional fragment of said E4 gene comprises the open reading frame 6 of E4.

8. The method according to claim 1 wherein said recombinant adeno-associated virus further comprises a second selected gene operatively linked to regulatory sequences capable of directing expression of said second gene, said second gene capable of facilitating the conversion of said single stranded recombinant virus to its double stranded form upon expression, and capable of co-expression with said first selected gene in the target cell, whereby the second gene product is expressed in the target cell, wherein said conversion occurs in said target cell, resulting in enhanced transduction of said recombinant virus in said target cell.

9. The method according to claim 8 wherein the regulatory sequences directing expression of said second gene comprise an inducible promoter, and wherein expression of said second gene occurs in the presence of an inducing agent.

10. A recombinant adeno-associated virus comprising (a) the DNA of at least a portion of the genome of an adeno-associated virus which portion is capable of transducing at least two selected genes or functional fragments thereof into a target cell in the absence of cell division; (b) a first selected gene operatively linked to regulatory sequences directing its expression, (c) a second selected gene operatively linked to regulatory sequences capable of directing expression of said second gene, said second gene capable of facilitating the conversion of said single stranded recombinant virus to its double stranded form upon expression, said first and said second genes flanked by the DNA of (a), said first and second gene capable of co-expression in the target cell.

11. The recombinant virus according to claim 10 wherein said second gene is selected from the group consisting of an adenovirus E4 gene, ORF6 of E4 and a functional fragment thereof.

12. The recombinant virus according to claim 10 wherein said second gene is the adenovirus E1 gene or a functional fragment thereof.

13. The recombinant virus according to claim 10 wherein the regulatory sequences directing expression of said second gene comprise an inducible promoter, and wherein expression of said second gene occurs in the presence of an inducing agent.



14. The recombinant virus according to claim 10 further comprising (d) an additional selected gene operatively linked to regulatory sequences capable of directing expression of said gene, said additional gene and said second gene capable of jointly facilitating the conversion of said single stranded recombinant virus to its double stranded form upon expression of both said second and additional genes, said first, second and additional genes flanked by the DNA of (a), and capable of co-expression in the target cell.

15. The recombinant virus according to claim 14 wherein said second gene is the adenovirus E4 gene or a functional fragment thereof and said additional gene is the adenovirus E1 gene or a functional fragment thereof.

16. The recombinant virus according to claim 15 wherein said functional fragment of E4 is the ORF6 sequence.

17. The recombinant virus according to claim 15 wherein the regulatory sequences directing expression of said additional gene comprise an inducible promoter, and wherein expression of said additional gene occurs in the presence of an inducing agent.

18. A method for enhancing the efficiency of transduction of a recombinant AAV into a target cell comprising the steps of: providing a recombinant adeno-associated virus of claim 10 through 16; infecting said target cells with said recombinant virus; and culturing said infected cells under conditions which enable expression of said selected genes in said target cell,

wherein said conversion occurs in said target cell, resulting in enhanced transduction of said recombinant virus in said target cell.

19. A method for enhancing the efficiency of transduction of a recombinant AAV into a target cell comprising the steps of: providing a recombinant adeno-associated virus of claim 17; infecting said target cells with said recombinant virus; and contacting said target cell with an inducing agent which induces the expression of said second or additional gene in said target cell, wherein said conversion occurs in said target cell, resulting in enhanced transduction of said recombinant virus in said target cell.

20. A pharmaceutical composition comprising a recombinant adeno-associated virus and an agent which facilitates the conversion of said ss recombinant virus to its double stranded form in a target cell.

21. The composition according to claim 20 wherein said virus is selected from the group consisting of the recombinant viruses of claims 10 - 16.

22. The composition according to claim 20 wherein said virus is the recombinant virus of claim 17 and further comprising an inducing agent which induces the expression of said second or additional gene in said recombinant virus.

23. A mammalian cell transduced with the recombinant adeno-associated viruses of claims 10-17.

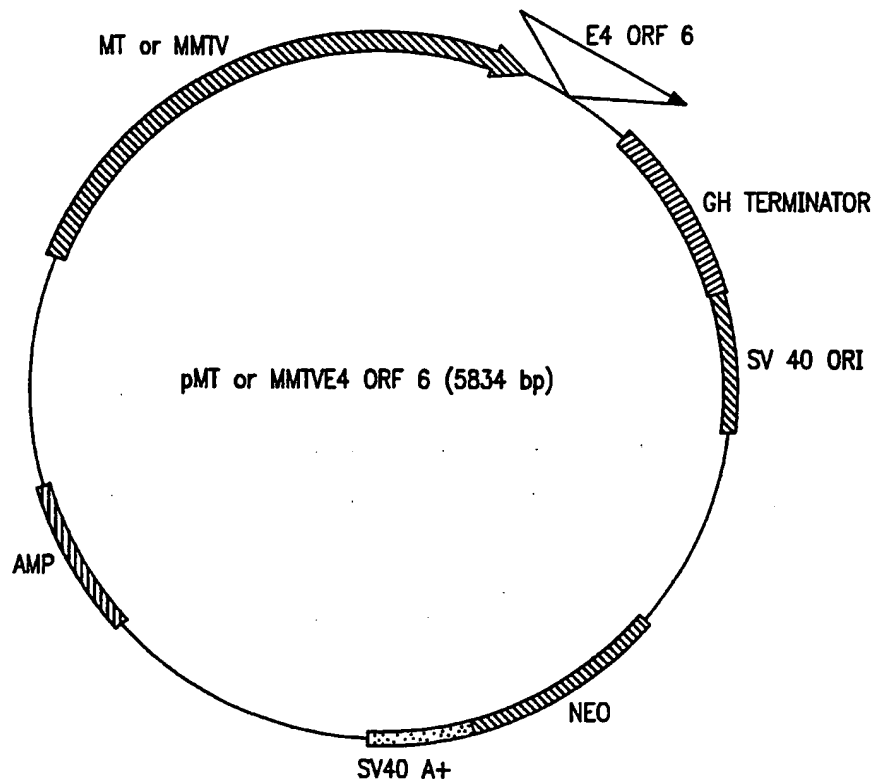
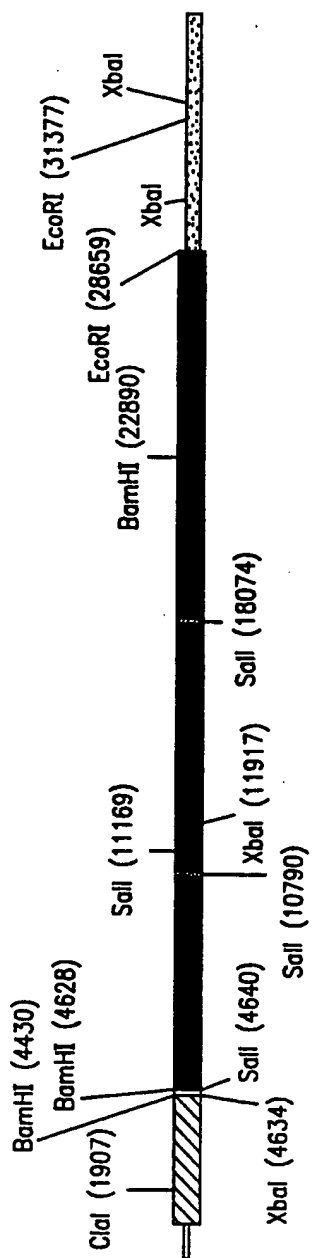


FIG. 1

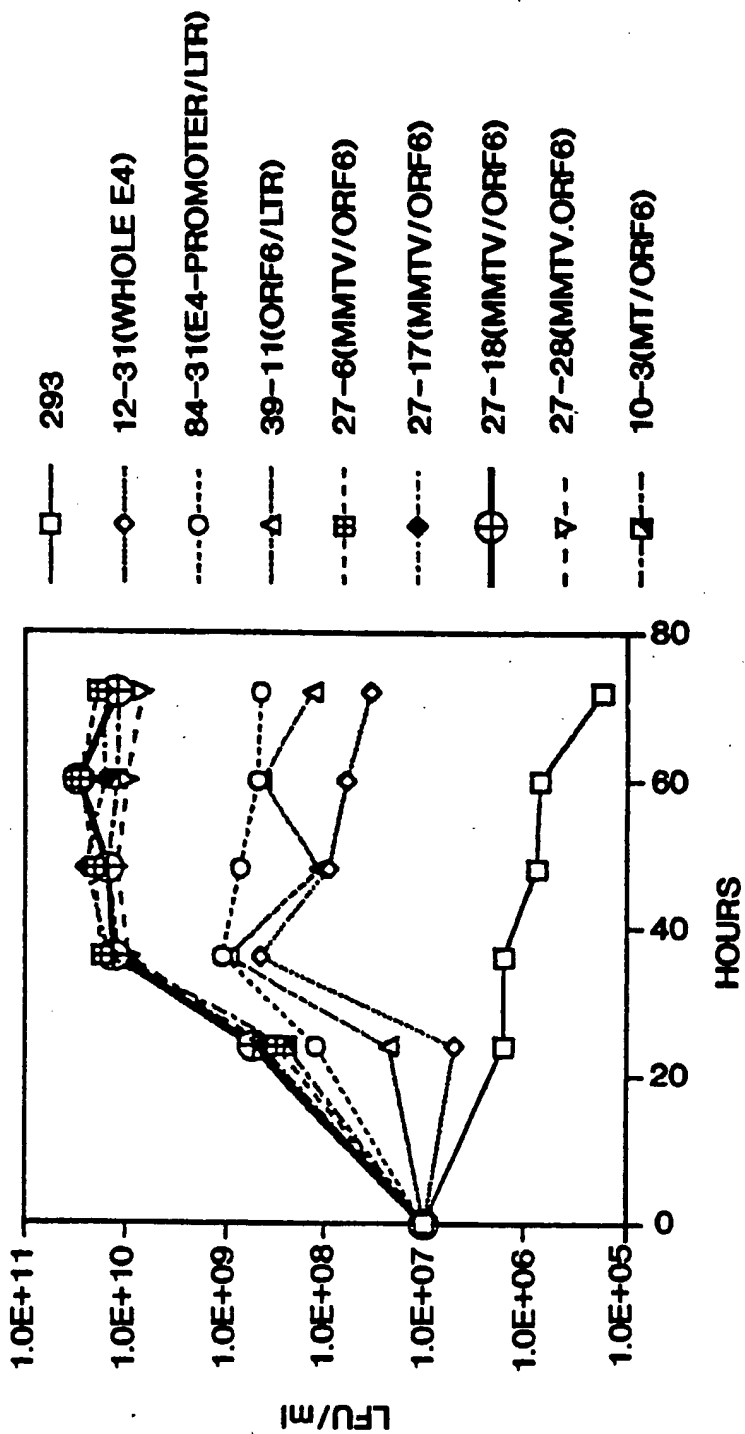


H5.001CB1acZ  
(35408 bp)

FIG. 2

3 / 11

FIG. 3



4 / 11

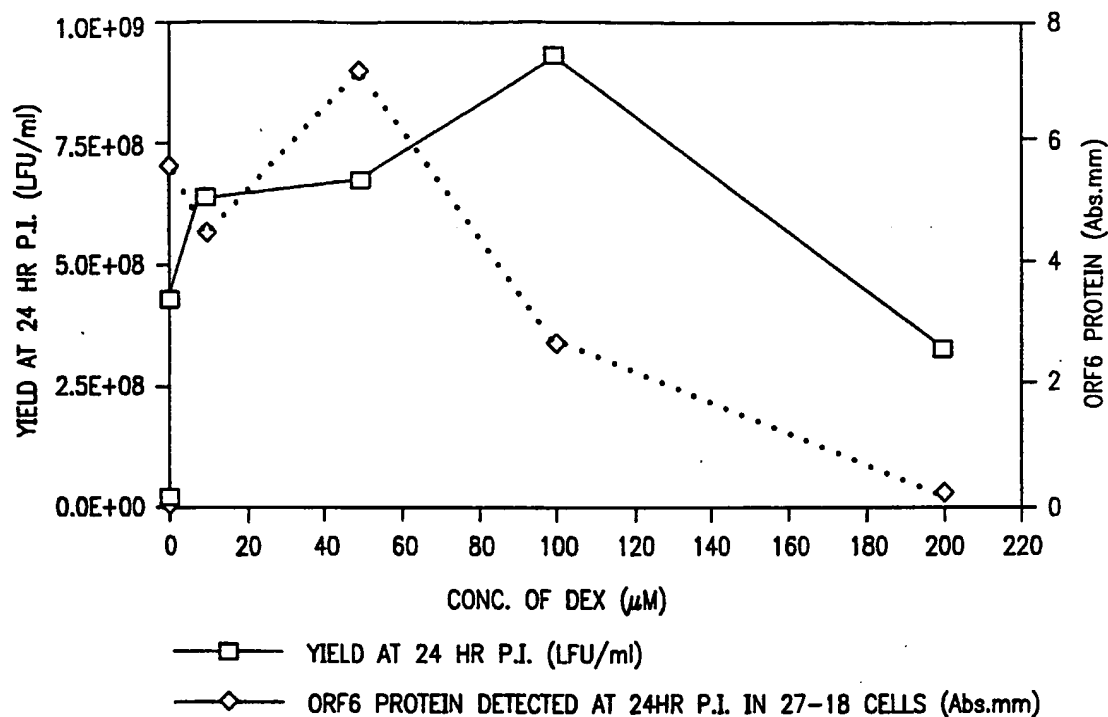


FIG. 4A

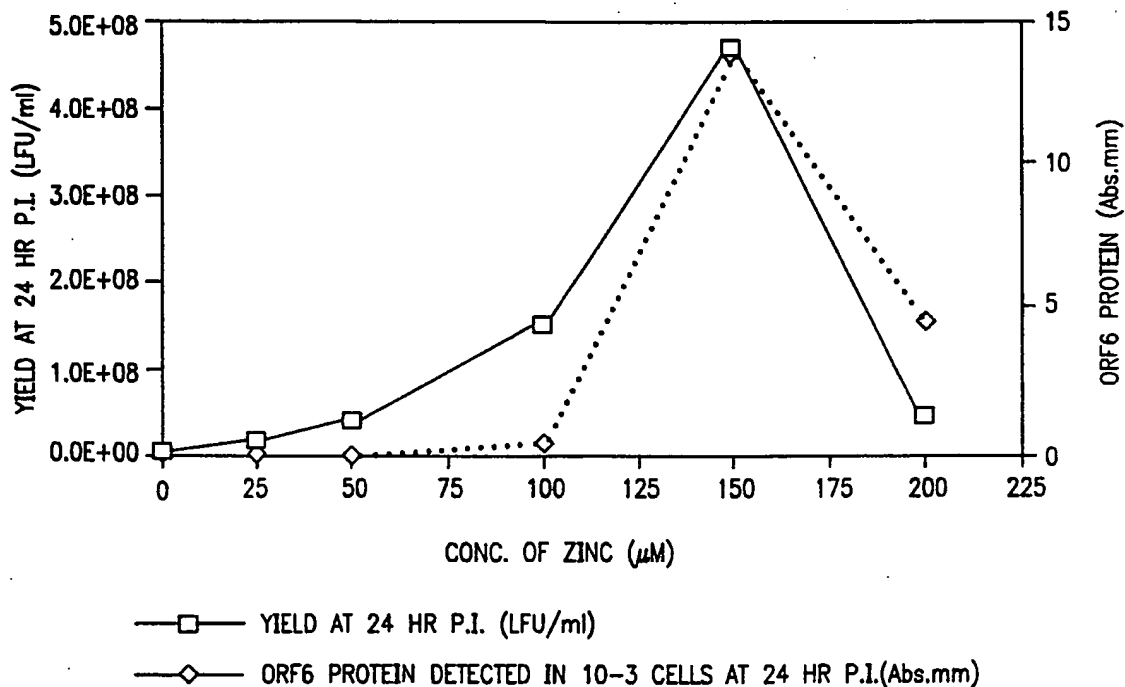


FIG. 4B

5/11

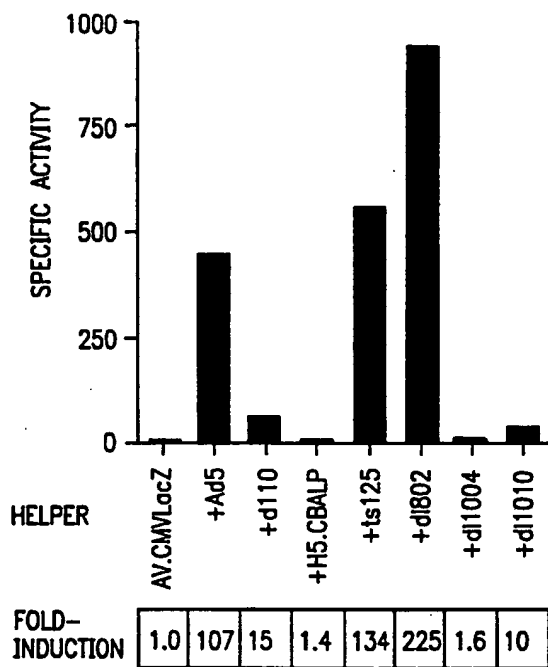


FIG. 5A

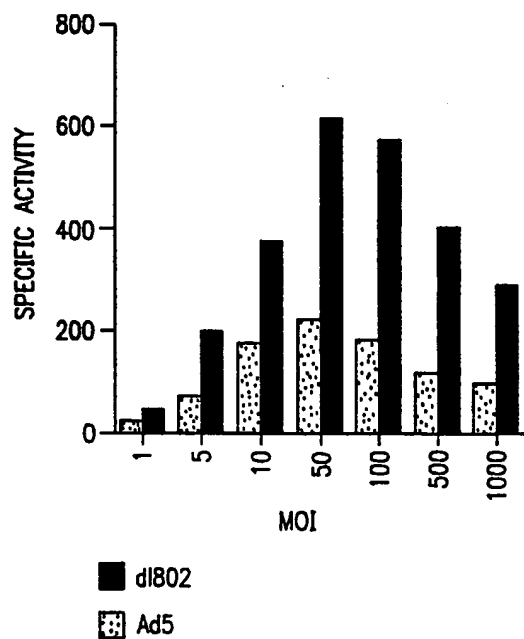


FIG. 5B

6/11

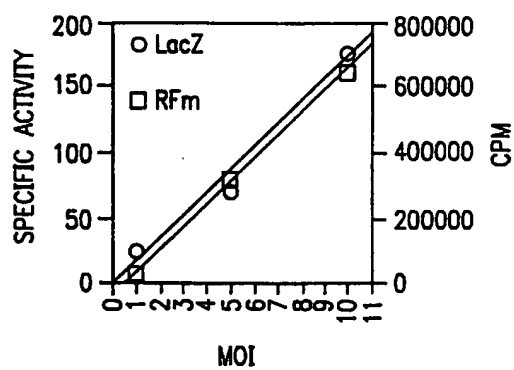


FIG. 6A

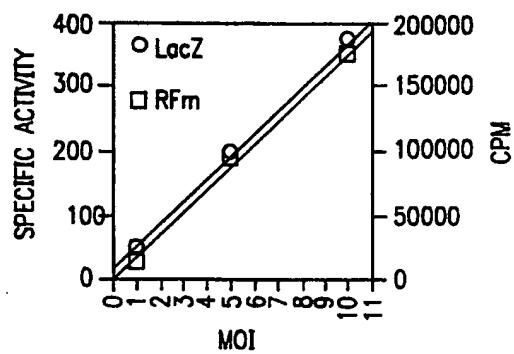


FIG. 6B



7/11

FIG. 7A

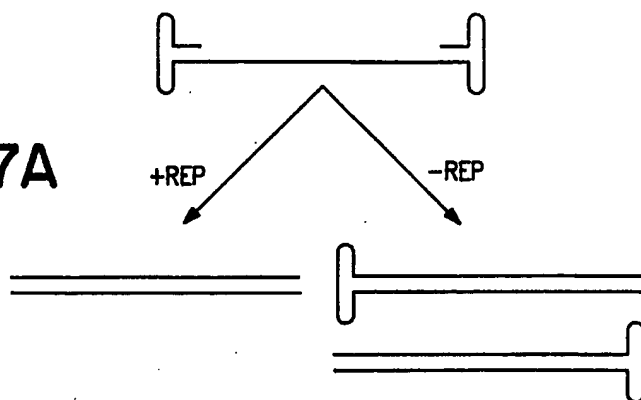


FIG. 7B

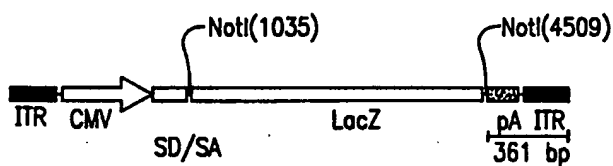
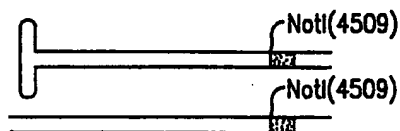


FIG. 7C



NotI DIGEST

FIG. 7D

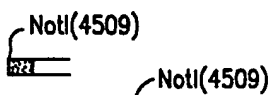
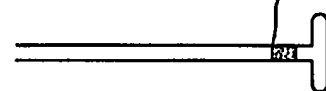


FIG. 7E



NotI DIGEST

FIG. 7F

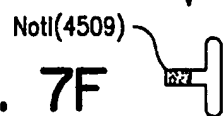


FIG. 8A

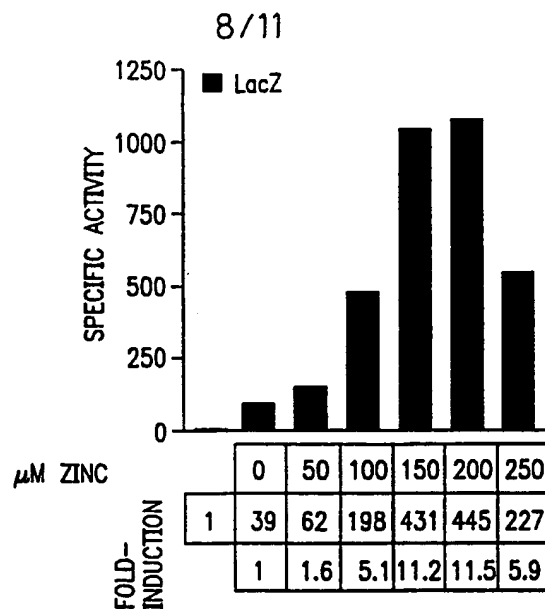


FIG. 8B

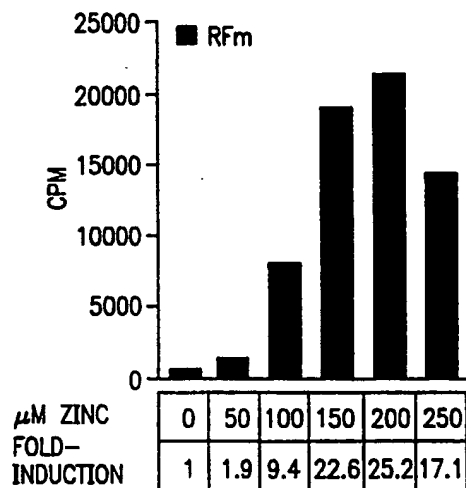
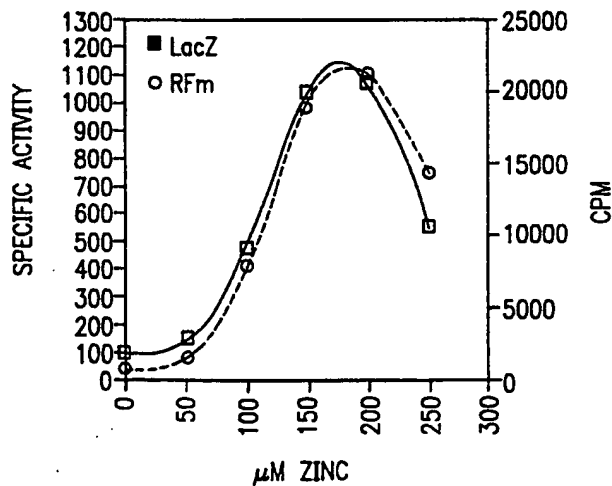


FIG. 8C



9 / 11

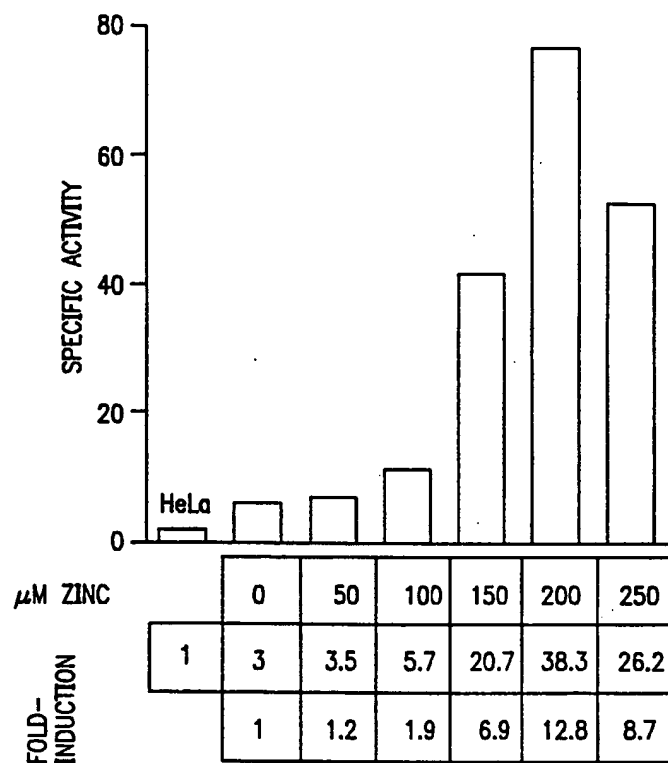


FIG. 9

10/11

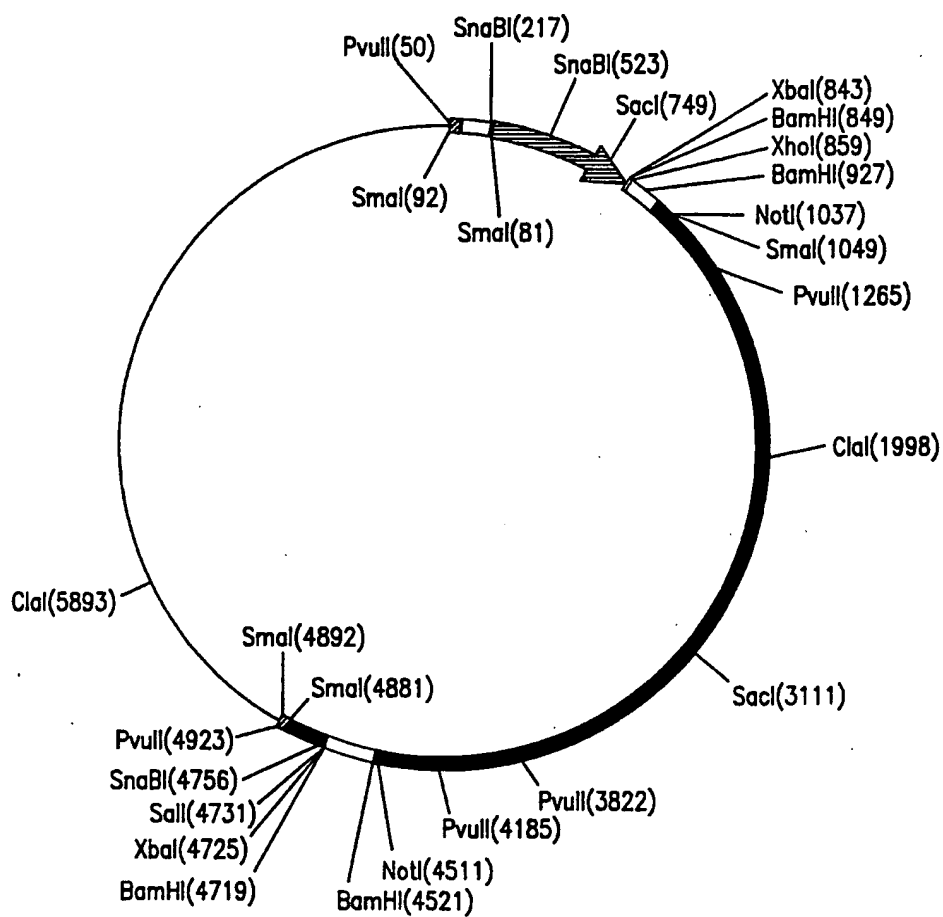


FIG. 10

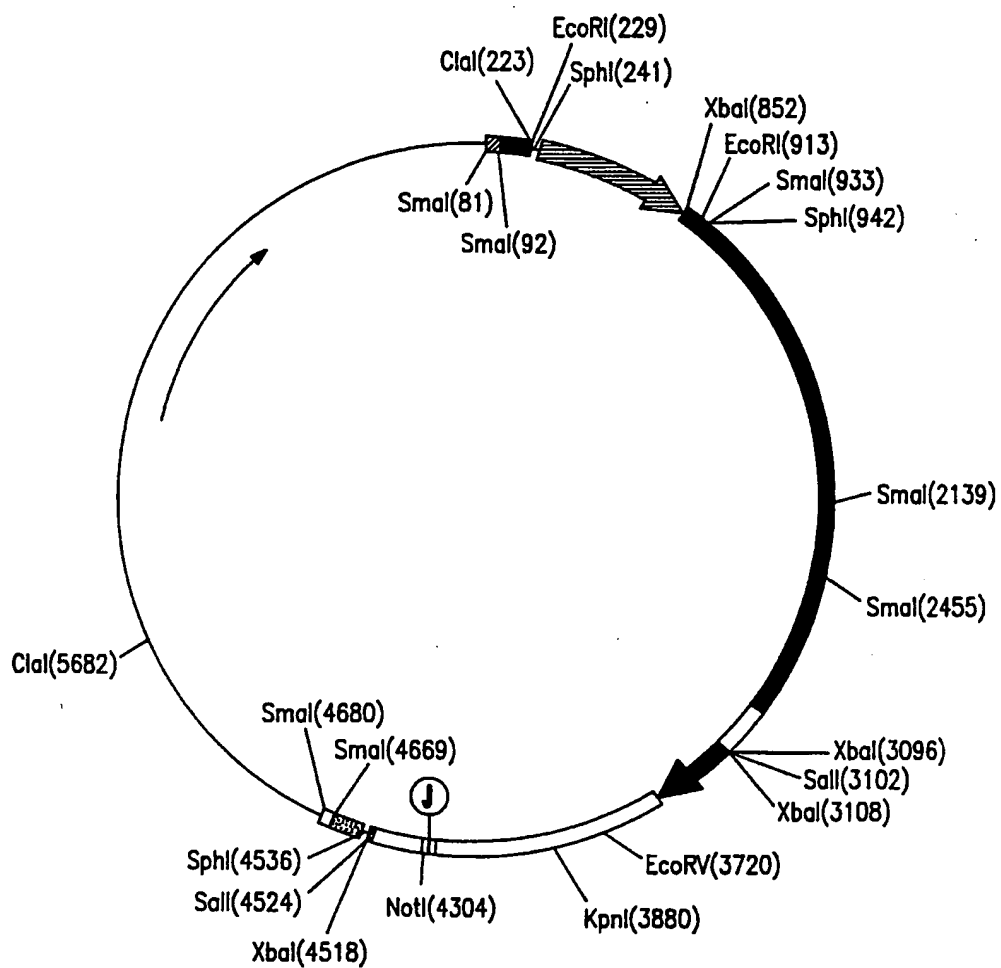


FIG. 11



## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

<b>(51) International Patent Classification <sup>6</sup>:</b> <b>C12N 15/86, 15/87, 5/10, A61K 35/76, 48/00</b>	<b>A3</b>	<b>(11) International Publication Number:</b> <b>WO 96/39530</b> <b>(43) International Publication Date:</b> 12 December 1996 (12.12.96)
<b>(21) International Application Number:</b> PCT/US96/10245 <b>(22) International Filing Date:</b> 4 June 1996 (04.06.96)  <b>(30) Priority Data:</b> 08/462,014      5 June 1995 (05.06.95)      US 08/549,489      27 October 1995 (27.10.95)      US  <b>(60) Parent Applications or Grants</b> <b>(63) Related by Continuation</b> US      08/462,014 (CIP) Filed on      5 June 1995 (05.06.95) US      08/549,489 (CIP) Filed on      27 October 1995 (27.10.95)  <b>(71) Applicant (for all designated States except US):</b> THE TRUSTEES OF THE UNIVERSITY OF PENNSYLVANIA [US/US]; 133 South 36th Street, Philadelphia, PA 19104-3246 (US).  <b>(72) Inventors; and</b> <b>(75) Inventors/Applicants (for US only):</b> WILSON, James, M. [US/US]; 1350 N. Avignon Drive, Gladwyne, PA 19035 (US). FISHER, Krishna, J. [US/US]; 4006 Pine Street, Philadelphia, PA 19104 (US). GAO, Guang-Ping [CN/US];		Robindale Apartment F5, 1925 Lawrence Road, Havertown, PA 19083 (US).  <b>(74) Agents:</b> BAK, Mary, E. et al.; Howson and Howson, Spring House Corporate Center, P.O. Box 457, Spring House, PA 19477 (US).  <b>(81) Designated States:</b> AL, AM, AU, AZ, BB, BG, BR, BY, CA, CN, CZ, EE, FI, GE, HU, IL, IS, JP, KE, KG, KP, KR, KZ, LK, LR, LS, LT, LV, MD, MG, MK, MN, MW, MX, NO, NZ, PL, RO, RU, SD, SG, SI, SK, TJ, TM, TR, TT, UA, UG, US, UZ, VN, ARIPO patent (KE, LS, MW, SD, SZ, UG), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, ML, MR, NE, SN, TD, TG).  <b>Published</b> <i>With international search report.</i>  <b>(88) Date of publication of the international search report:</b> 5 June 1997 (05.06.97)
<b>(54) Title:</b> RECOMBIANT ADENOVIRUS AND ADENO-ASSOCIATED VIRUS, CELL LINES, AND METHODS OF PRODUCTION AND USE THEREOF		
<b>(57) Abstract</b> <p>An adenovirus E1/E4 expressing packaging cell line is provided, which permits the generation of recombinant adenoviruses deleted in both gene regions. A method for enhancing the efficiency of transduction of a recombinant AAV into a target cell is provided by infecting a target cell with a recombinant AAV comprising a selected transgene under the control of regulatory sequences. The infected cell is contacted with an agent which facilitates the conversion of single stranded recombinant virus to its double stranded form.</p>		

**FOR THE PURPOSES OF INFORMATION ONLY**

Codes used to identify States party to the PCT on the front pages of pamphlets publishing international applications under the PCT.

AM	Armenia	GB	United Kingdom	MW	Malawi
AT	Austria	GE	Georgia	MX	Mexico
AU	Australia	GN	Guinea	NE	Niger
BB	Barbados	GR	Greece	NL	Netherlands
BE	Belgium	HU	Hungary	NO	Norway
BF	Burkina Faso	IE	Ireland	NZ	New Zealand
BG	Bulgaria	IT	Italy	PL	Poland
BJ	Benin	JP	Japan	PT	Portugal
BR	Brazil	KE	Kenya	RO	Romania
BY	Belarus	KG	Kyrgyzstan	RU	Russian Federation
CA	Canada	KP	Democratic People's Republic of Korea	SD	Sudan
CF	Central African Republic	KR	Republic of Korea	SE	Sweden
CG	Congo	KZ	Kazakhstan	SG	Singapore
CH	Switzerland	LI	Liechtenstein	SI	Slovenia
CI	Côte d'Ivoire	LK	Sri Lanka	SK	Slovakia
CM	Cameroon	LR	Liberia	SN	Senegal
CN	China	LT	Lithuania	SZ	Swaziland
CS	Czechoslovakia	LU	Luxembourg	TD	Chad
CZ	Czech Republic	LV	Latvia	TG	Togo
DE	Germany	MC	Monaco	TJ	Tajikistan
DK	Denmark	MD	Republic of Moldova	TT	Trinidad and Tobago
EE	Estonia	MG	Madagascar	UA	Ukraine
ES	Spain	ML	Mali	UG	Uganda
FI	Finland	MN	Mongolia	US	United States of America
FR	France	MR	Mauritania	UZ	Uzbekistan
GA	Gabon			VN	Viet Nam

# INTERNATIONAL SEARCH REPORT

International Application No.  
PCT/US 96/10245

## A. CLASSIFICATION OF SUBJECT MATTER

IPC 6 C12N15/86 C12N15/87 C12N5/10 A61K35/76 A61K48/00

According to International Patent Classification (IPC) or to both national classification and IPC

## B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 C12N A61K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

## C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P,X	J.VIROLOGY, vol. 70, no. 1, January 1996, pages 520-532, XP000570468 FISHER, K.J. ET AL.: "Transduction with recombinant Adeno-Associated Virus for gene therapy is limited by leading-strand synthesis"	1-7
P,Y	see the whole document ---	8-23
P,X	J.VIROLOGY, vol. 70, no. 3, March 1996, pages 1845-1854, XP002027273 WEITZMAN, M.D. ET AL.: "Recruitment of wild-type and recombinant Adeno-Associated Virus into Adenovirus replication centers"	1-7
P,Y	see the whole document ---	8-23
	-/--	

☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

### \* Special categories of cited documents:

- \*A\* document defining the general state of the art which is not considered to be of particular relevance
- \*E\* earlier document but published on or after the international filing date
- \*L\* document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- \*O\* document referring to an oral disclosure, use, exhibition or other means
- \*P\* document published prior to the international filing date but later than the priority date claimed

- \*T\* later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
- \*X\* document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
- \*Y\* document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
- \*&\* document member of the same patent family

Date of the actual completion of the international search

10 March 1997

Date of mailing of the international search report

25. 03. 97

Name and mailing address of the ISA

European Patent Office, P.B. 5818 Patentlaan 2  
NL - 2280 HV Rijswijk  
Tel. (+ 31-70) 340-2040, Tx. 31 651 epo nl,  
Fax: (+ 31-70) 340-3016

Authorized officer

Donath, C



# INTERNATIONAL SEARCH REPORT

Int. l. Application No.  
PCT/US 96/10245

## C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
P,X	<p>WO 95 20671 A (RHONE-POULENC RORER S.A.) 3 August 1995 see page 2, line 33 - page 3, line 30 see page 5, line 28 - page 7, line 24; examples 2-5</p> <p style="text-align: center;">---</p>	1-7
P,X	<p>WO 96 13598 A (THE TRUSTEES OF THE UNIVERSITY OF PENNSYLVANIA) 9 May 1996 see page 4, line 23 - page 5, line 25 see page 8, line 1 - page 15, line 13 see page 18, line 28 - page 19, line 13</p> <p style="text-align: center;">---</p>	1-7
E	<p>WO 96 22378 A (RHONE-POULENC RORER S.A.) 25 July 1996 see page 3, line 3 - line 21 see page 7, line 17 - line 32 see page 13, line 27 - page 16, line 26</p> <p style="text-align: center;">---</p>	1-7
X	<p>GENE, vol. 119, no. 2, 1992, pages 265-272, XP002027274 NAHREINI, P. ET AL.: "Cloning and integration of DNA fragments in human cells via the inverted terminal repeats of the adeno-associated virus 2 genome" see the whole document</p> <p style="text-align: center;">---</p>	1-7
Y	<p>CRC HANDBOOK OF PARVOVIRUSES, ED.P.TIJSSER, vol. 1, 1990, pages 155-168, XP002027275 CARTER, B.J.: "The growth cycle of adeno-associated virus" see page 158, last paragraph - page 161, paragraph 1</p> <p style="text-align: center;">-----</p>	8-23

# INTERNATIONAL SEARCH REPORT

Information on patent family members

International Application No

PCT/US 96/10245

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO 9520671 A	03-08-95	FR 2716682 A	01-09-95
		AU 1539595 A	15-08-95
		CA 2181602 A	03-08-95
		EP 0741793 A	13-11-96
		FI 962990 A	26-07-96
		NO 962950 A	12-07-96
		ZA 9500628 A	23-10-95
-----			
WO 9613598 A	09-05-96	AU 4405596 A	23-05-96
-----			
WO 9622378 A	25-07-96	FR 2729674 A	26-07-96
		AU 4544396 A	07-08-96
-----			